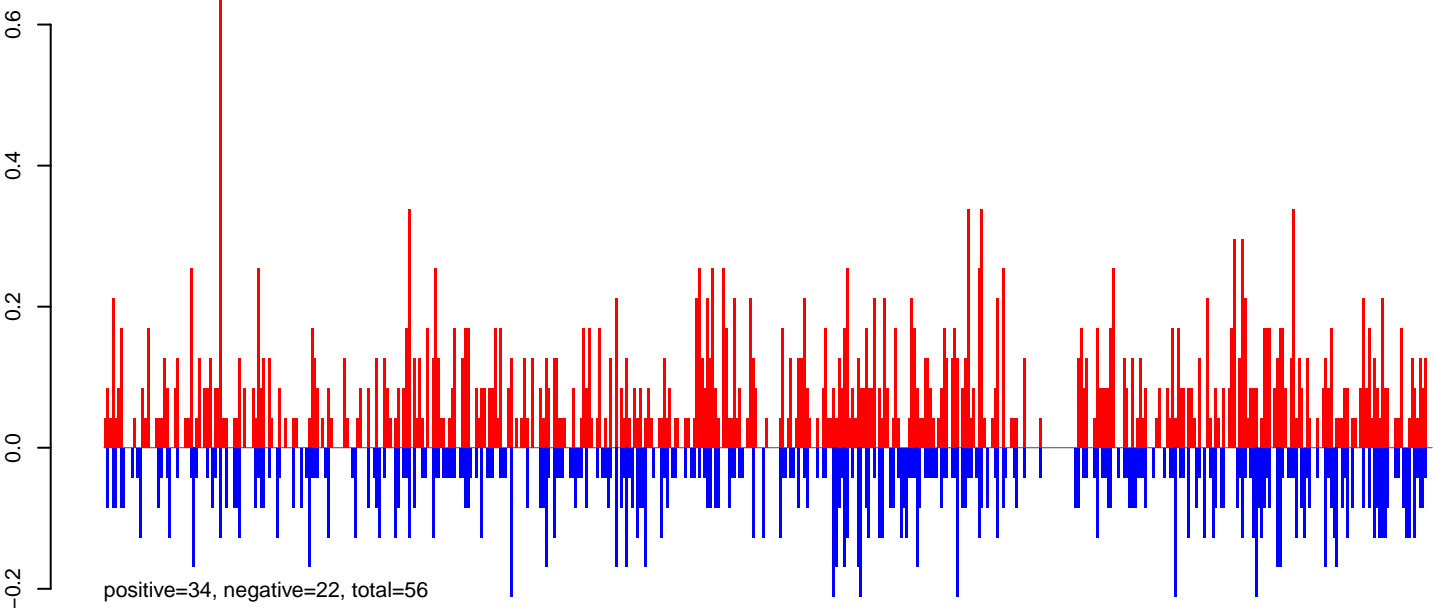
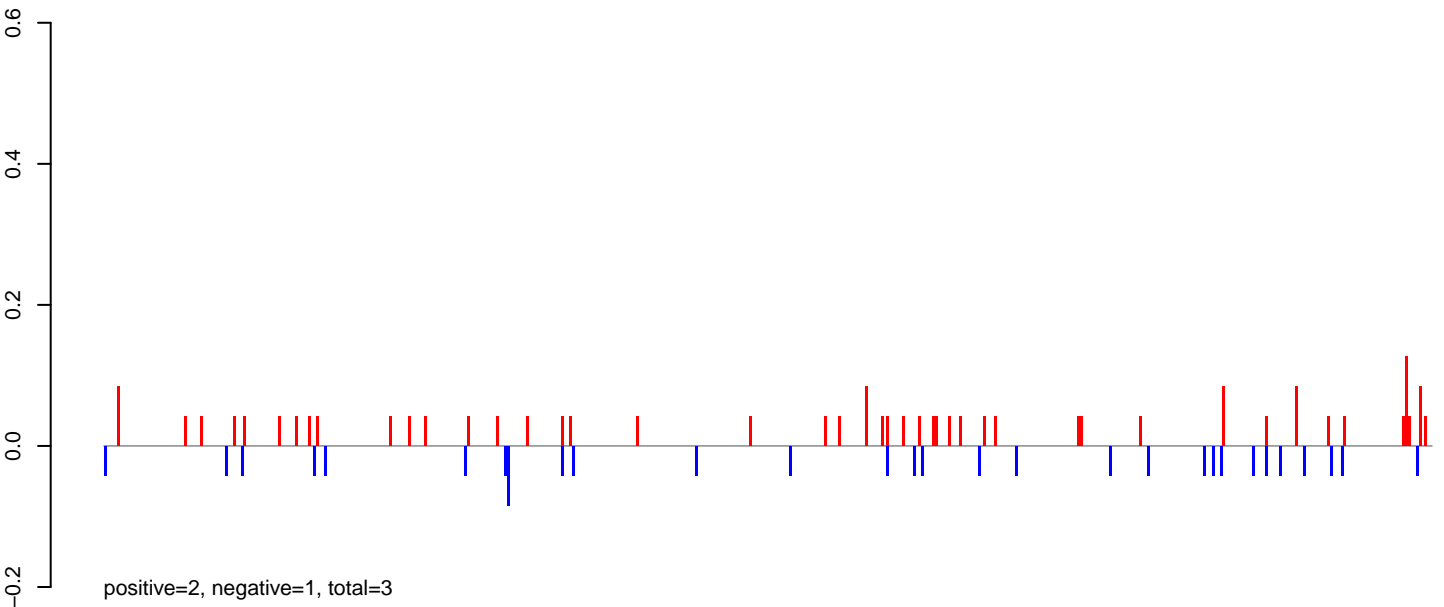


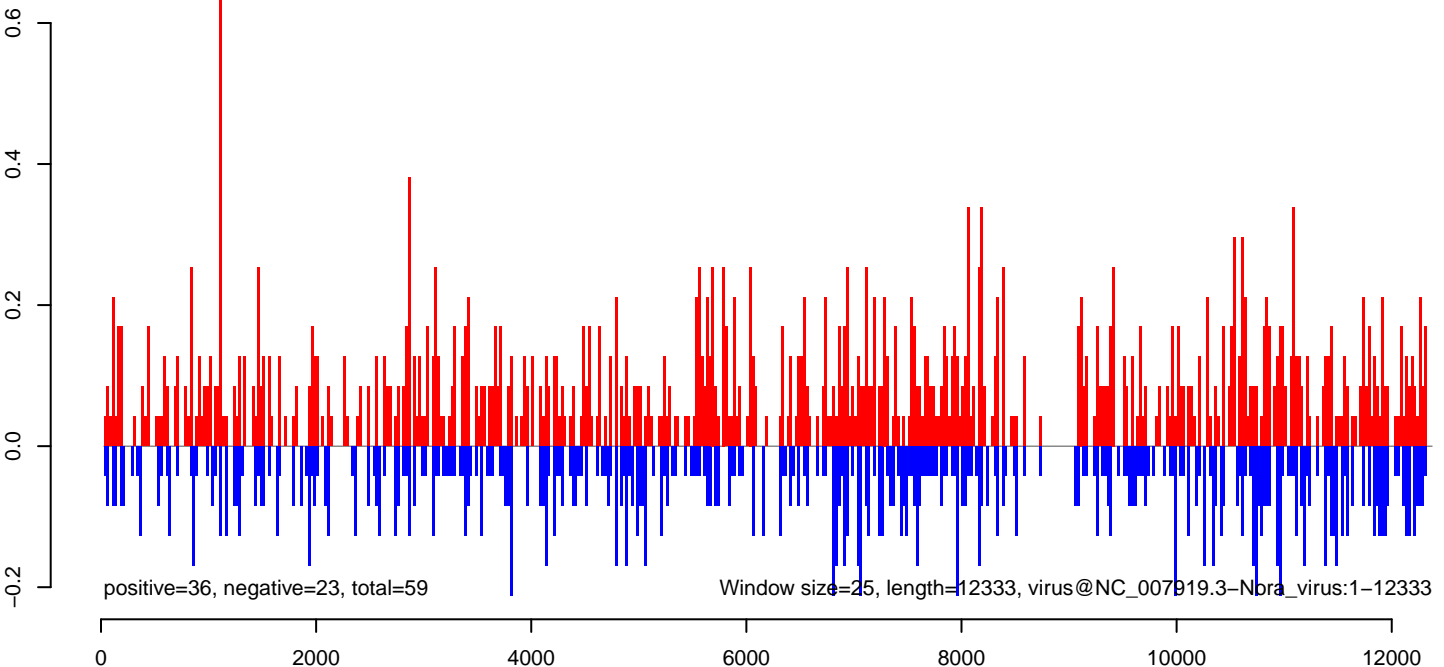
AeAlbo_C636_cells.18_23.rep



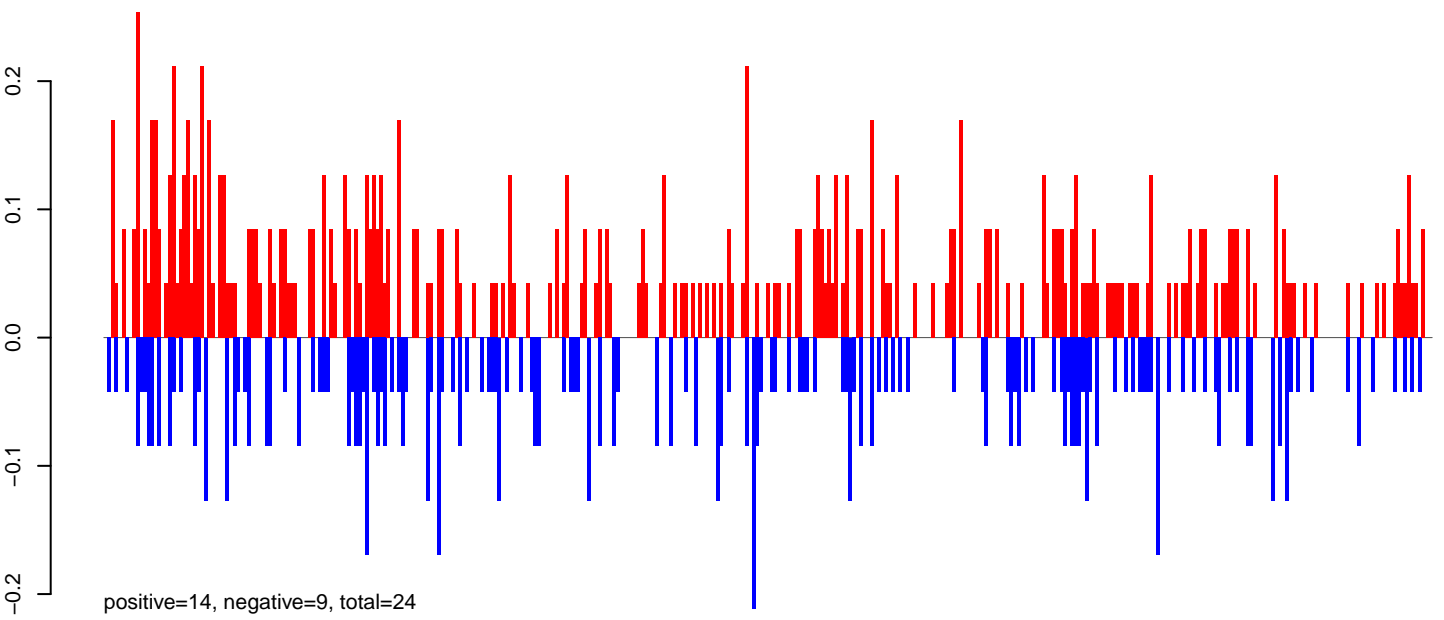
AeAlbo_C636_cells.24_35.rep



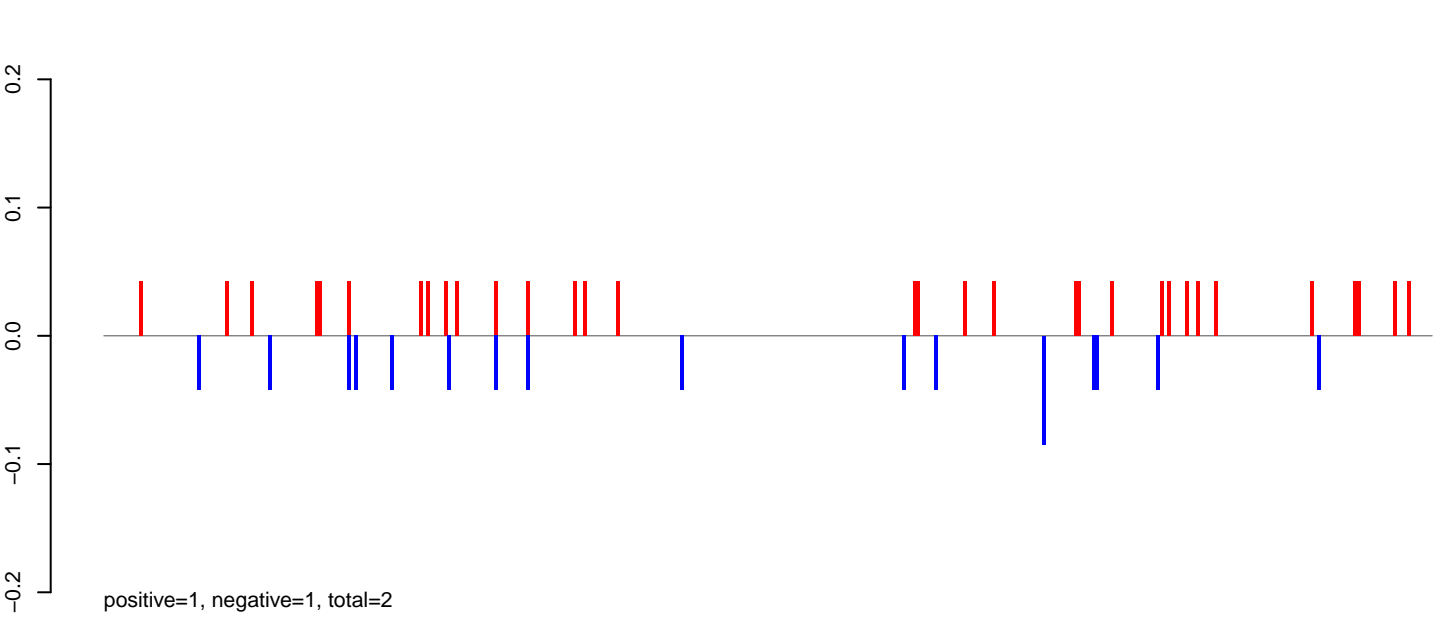
AeAlbo_C636_cells.rep



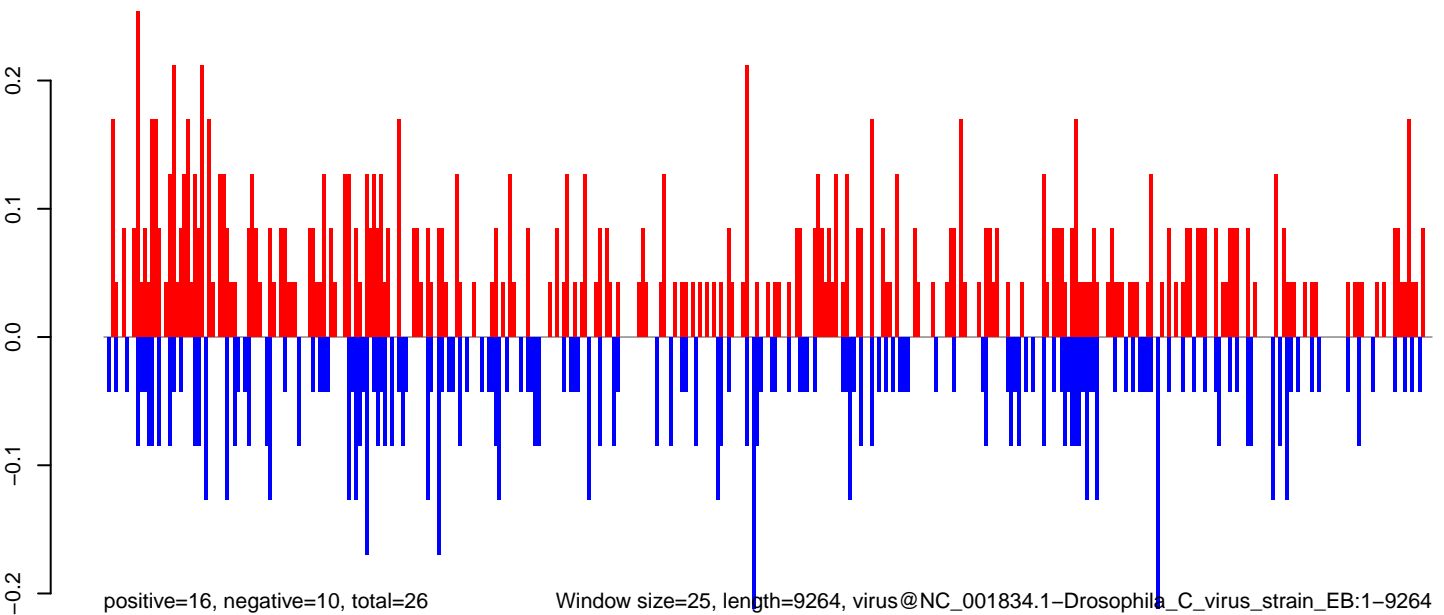
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

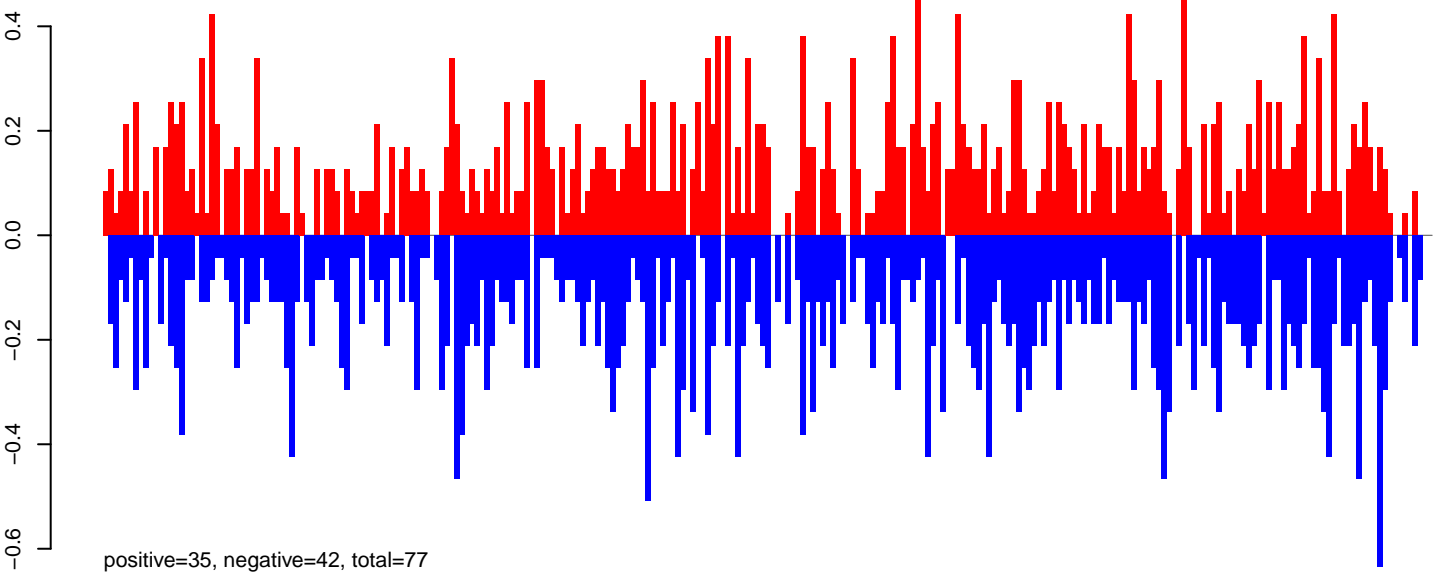


AeAlbo_C636_cells.rep

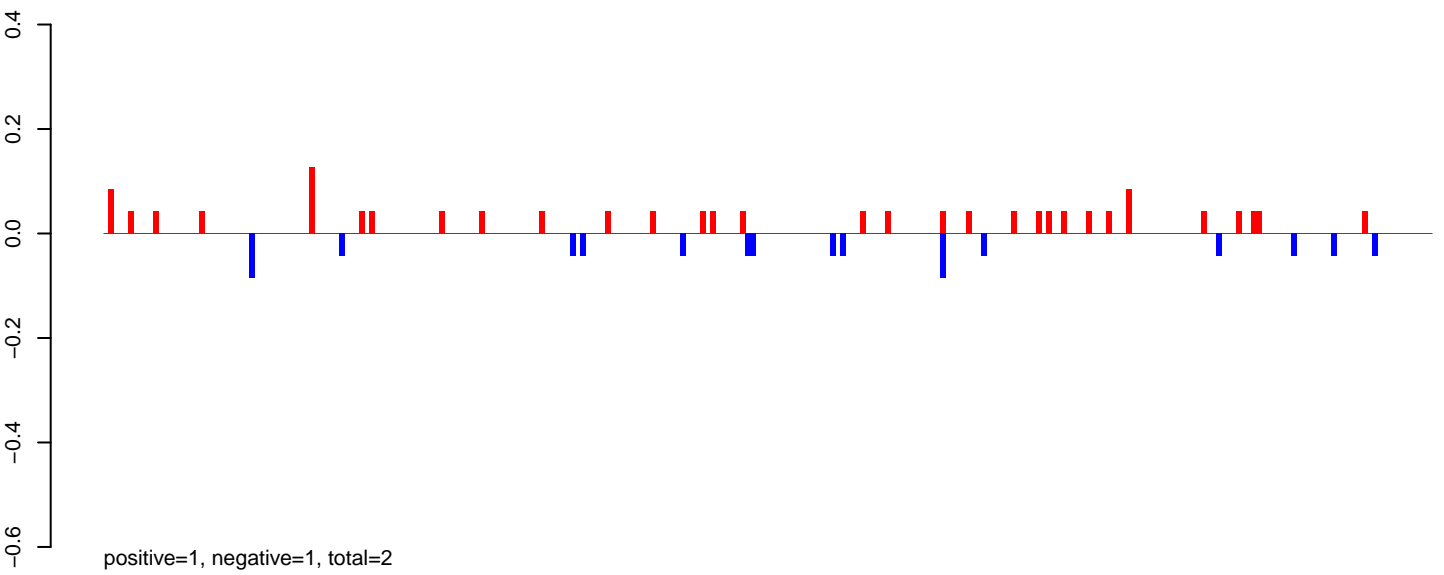


0 2000 4000 6000 8000

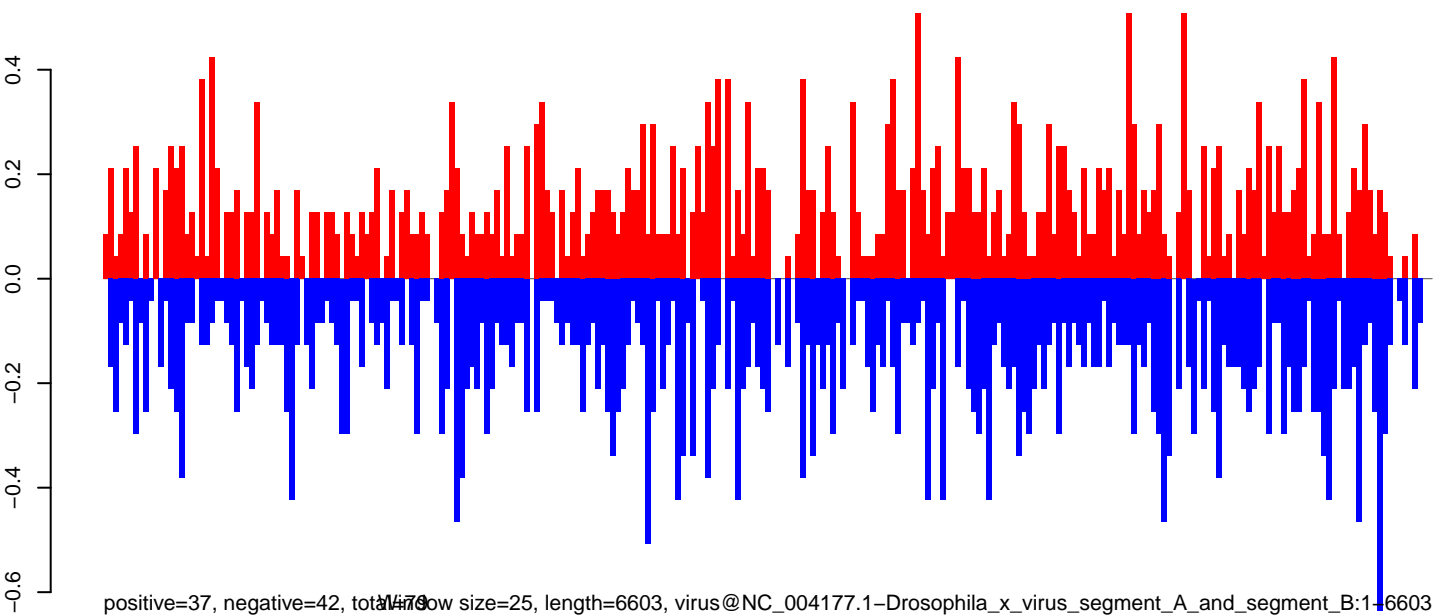
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

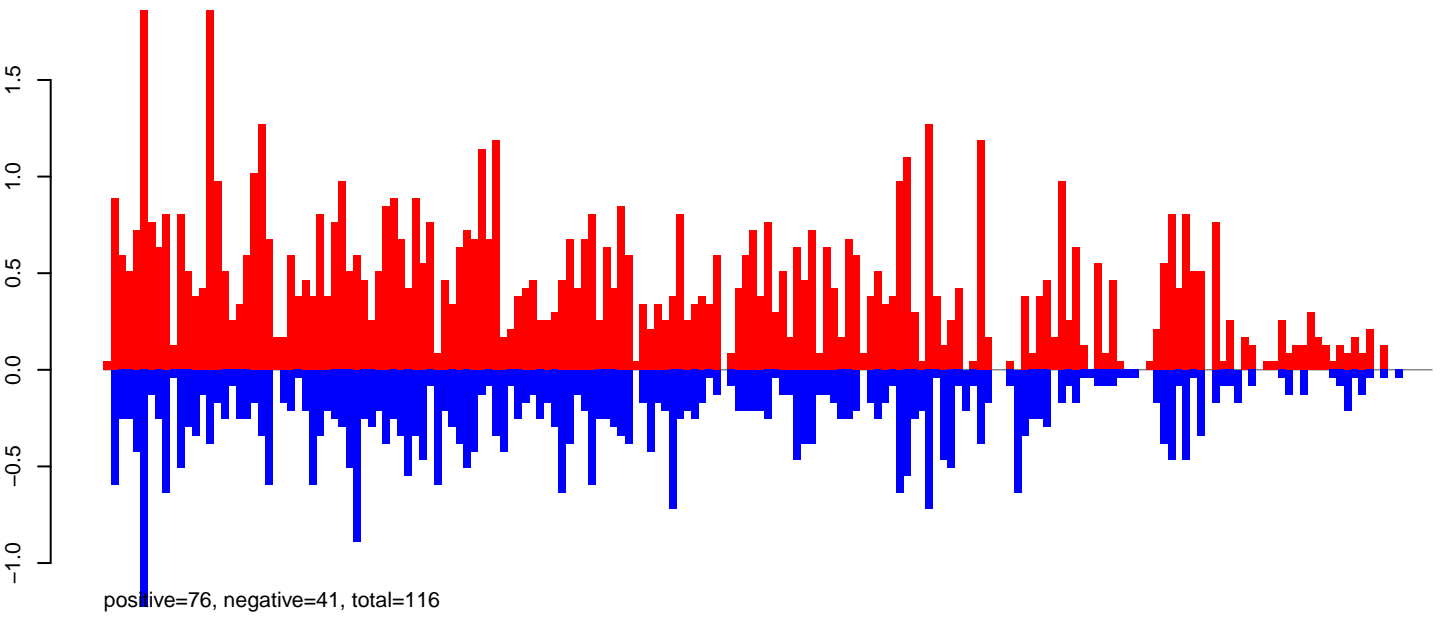


AeAlbo_C636_cells.rep

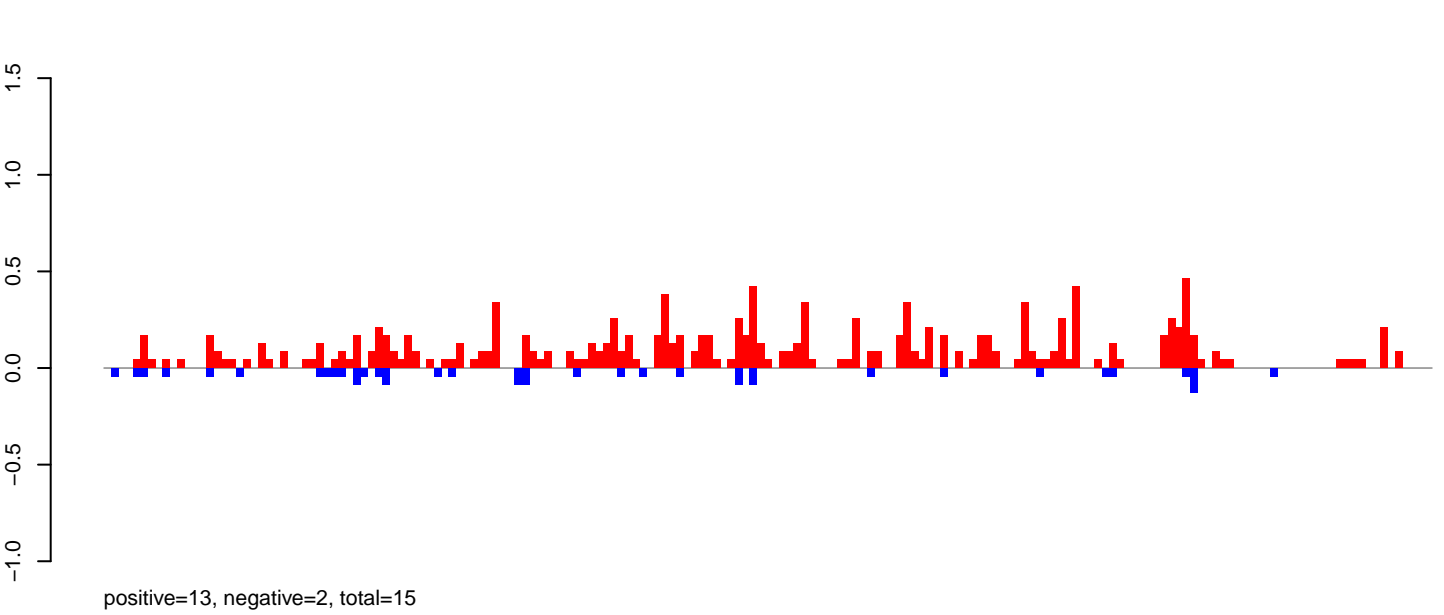


row size=25, length=6603, virus@NC_004177.1-Drosophila_x_virus_segment_A_and_segment_B:1-6603

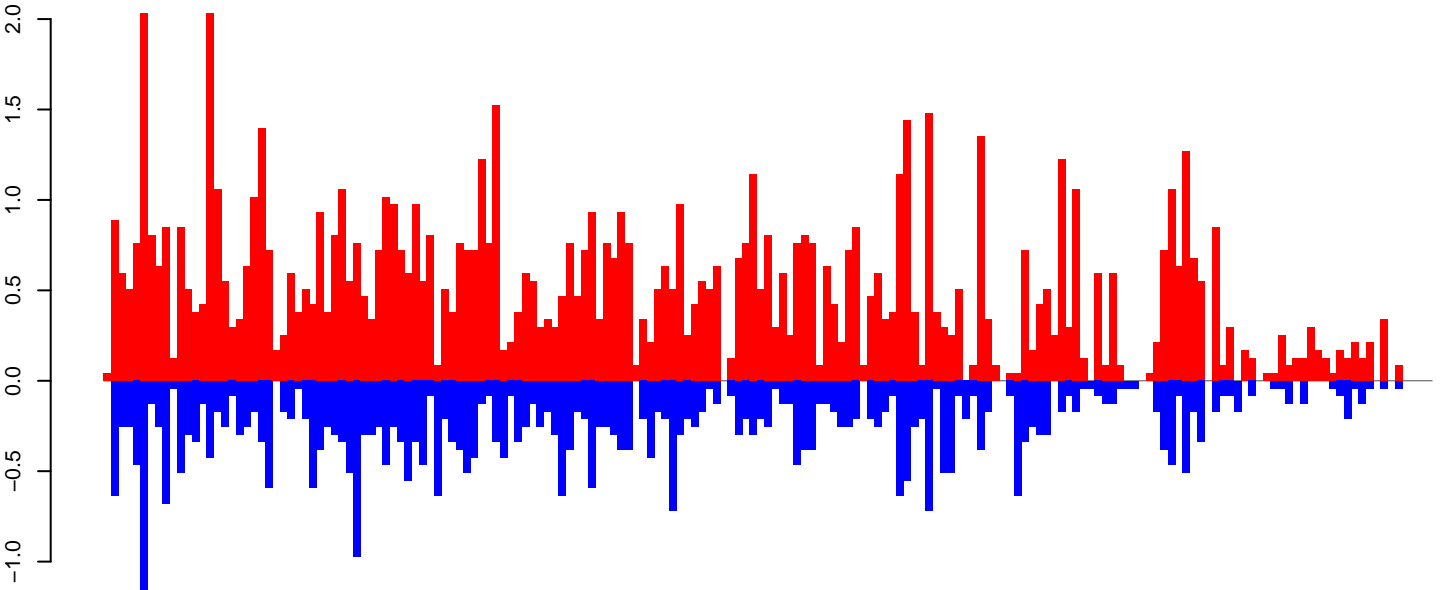
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



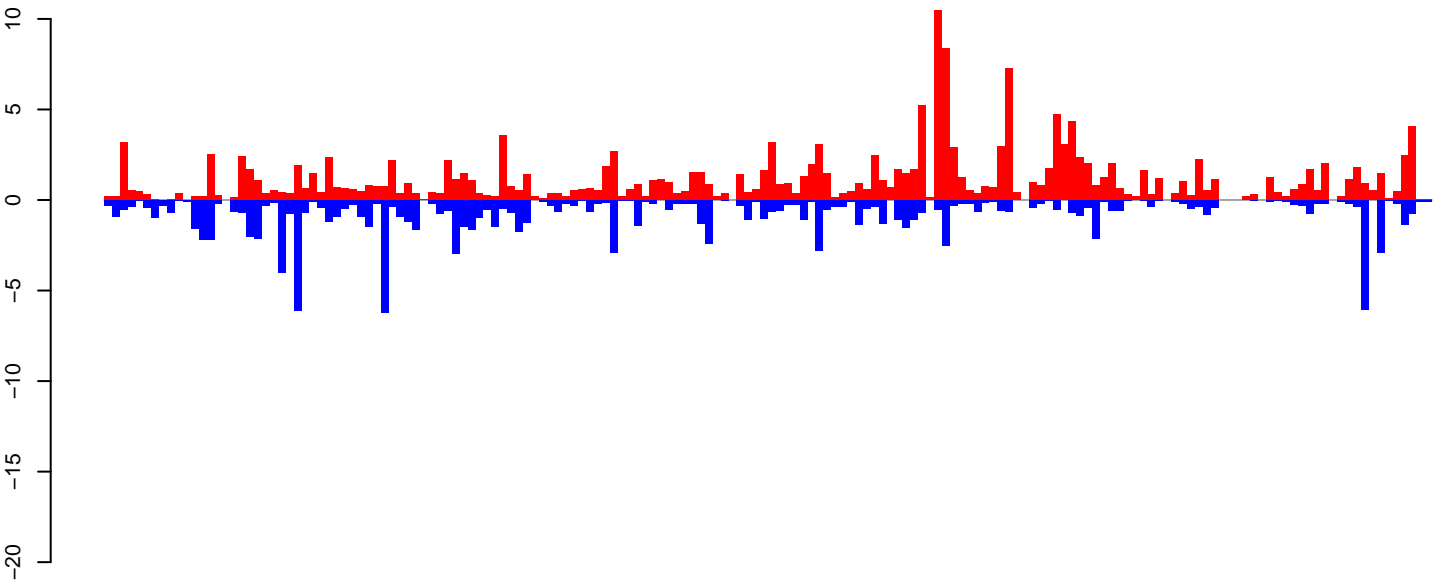
AeAlbo_C636_cells.rep



ze=25, col=95, row=42, @G812065.1-Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

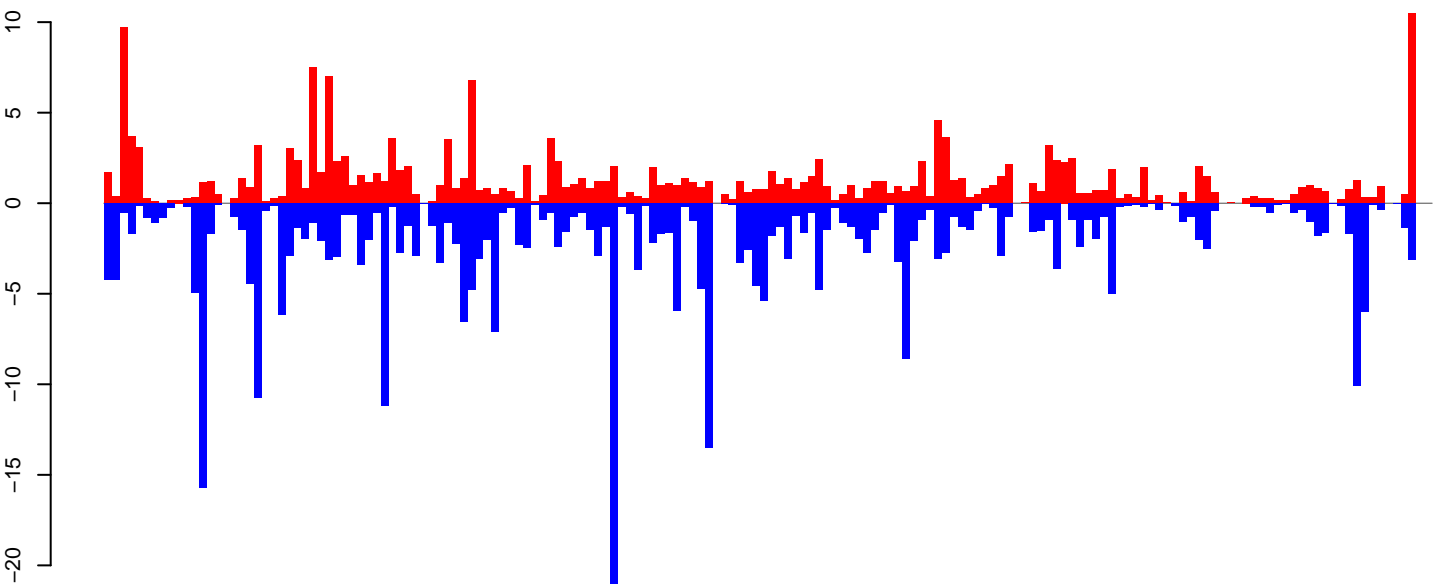
0 1000 2000 3000 4000

AeAlbo_C636_cells.18_23.rep



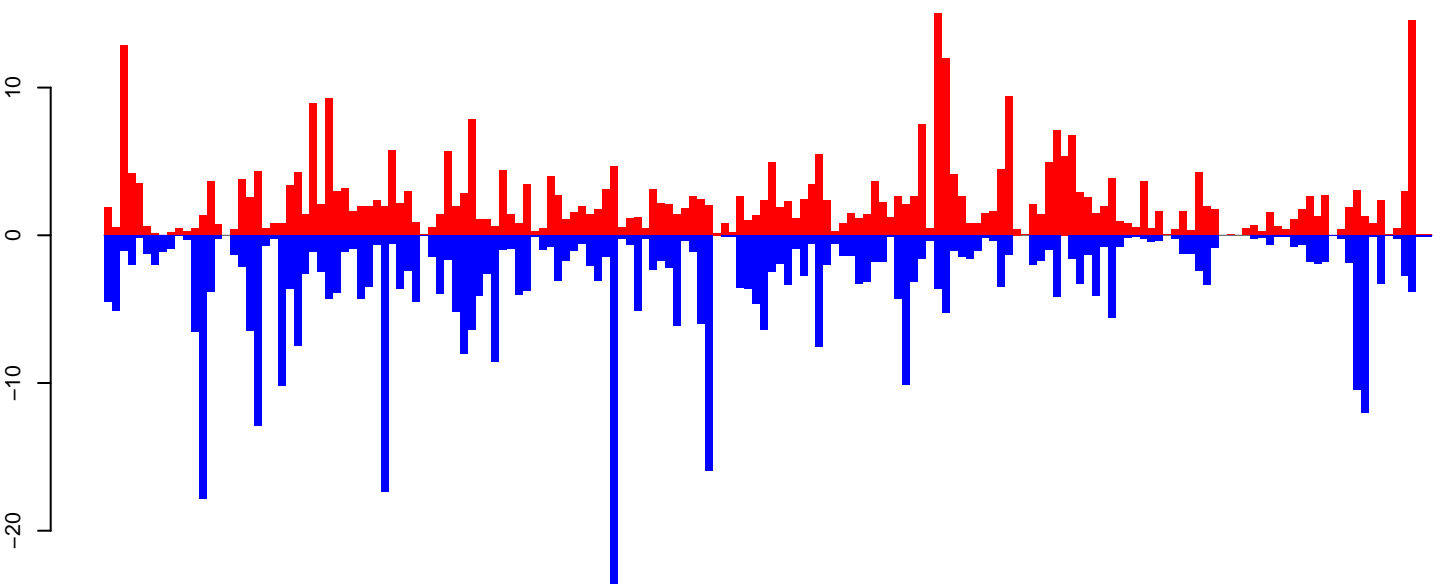
positive=188, negative=117, total=305

AeAlbo_C636_cells.24_35.rep



positive=205, negative=334, total=540

AeAlbo_C636_cells.rep

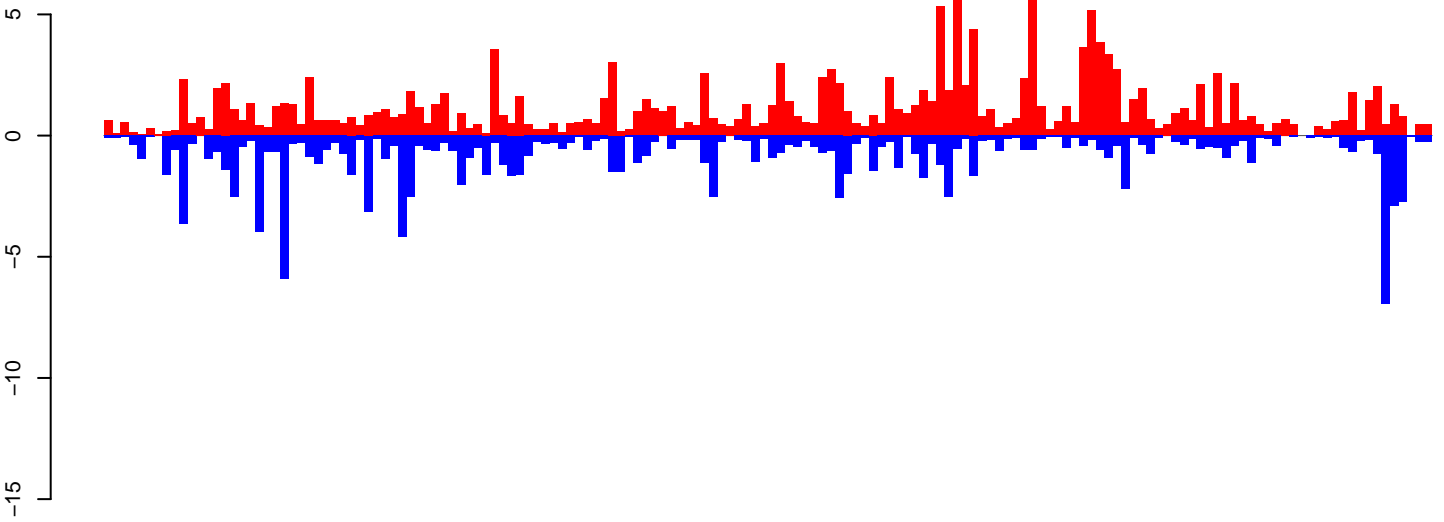


positive=393, negative=452, total=845

Window size=25, length=4176, virus@X74945.1-Aedes_albopictus_densovirus_2_DNA:1-4176

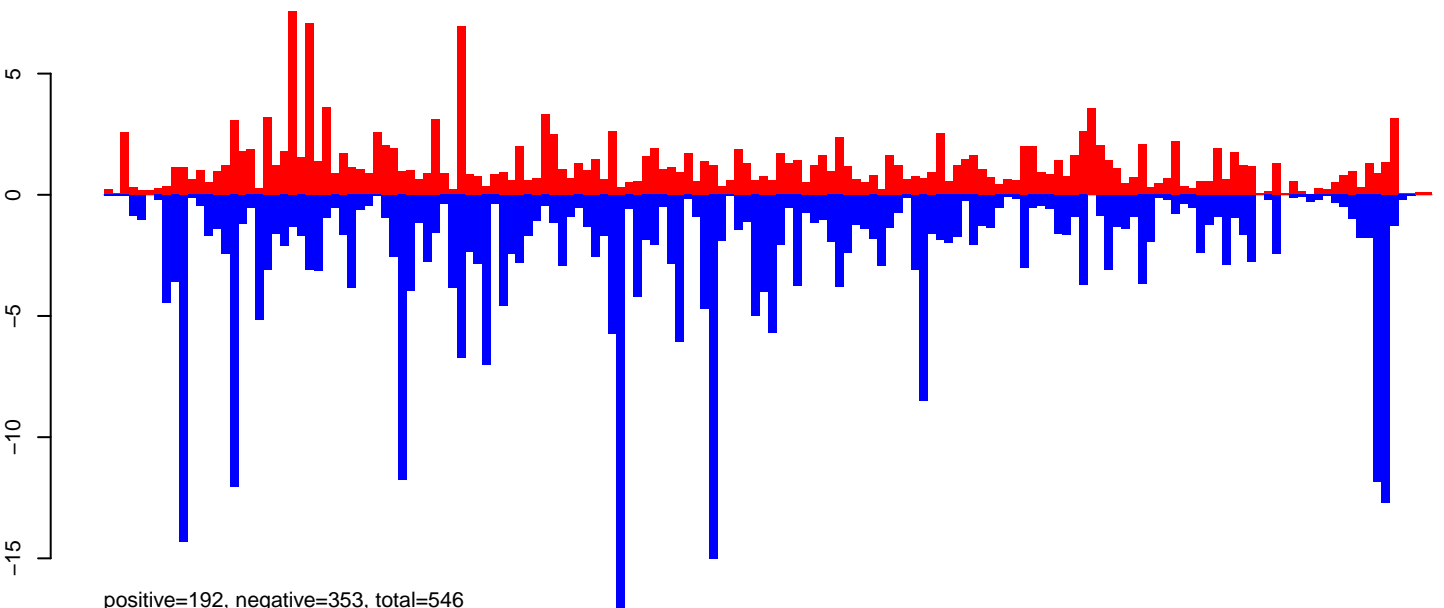
0 1000 2000 3000 4000

AeAlbo_C636_cells.18_23.rep



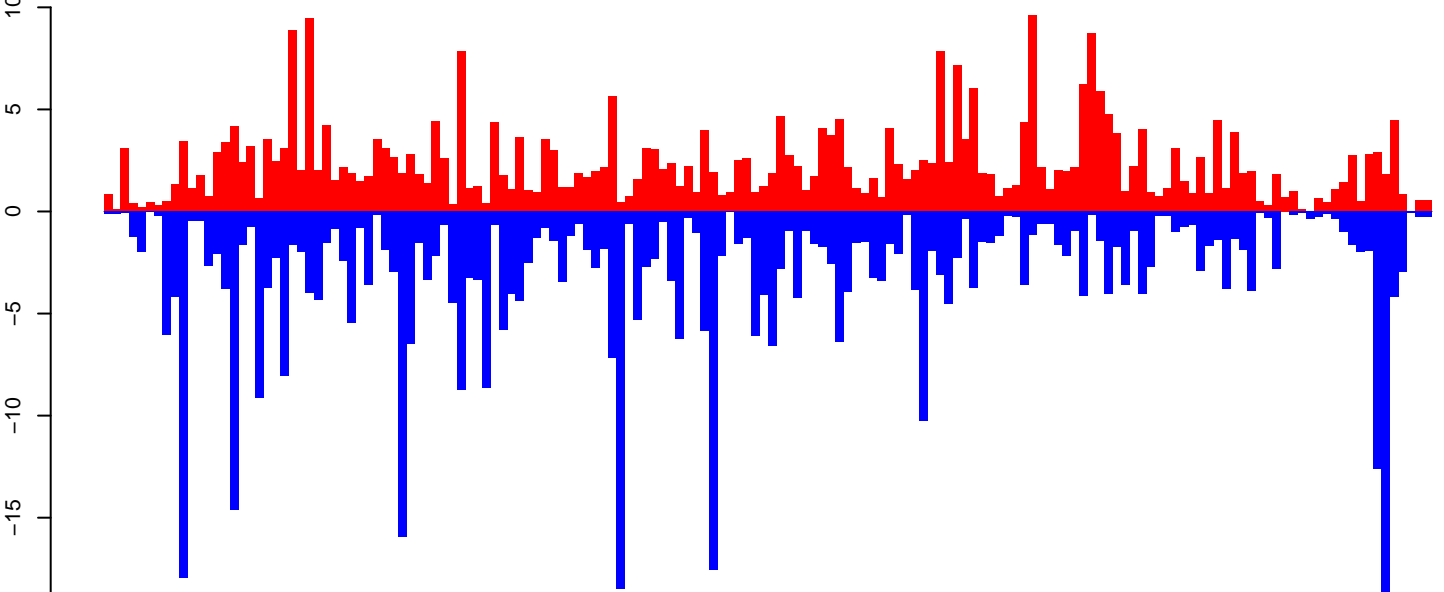
positive=180, negative=123, total=303

AeAlbo_C636_cells.24_35.rep



positive=192, negative=353, total=546

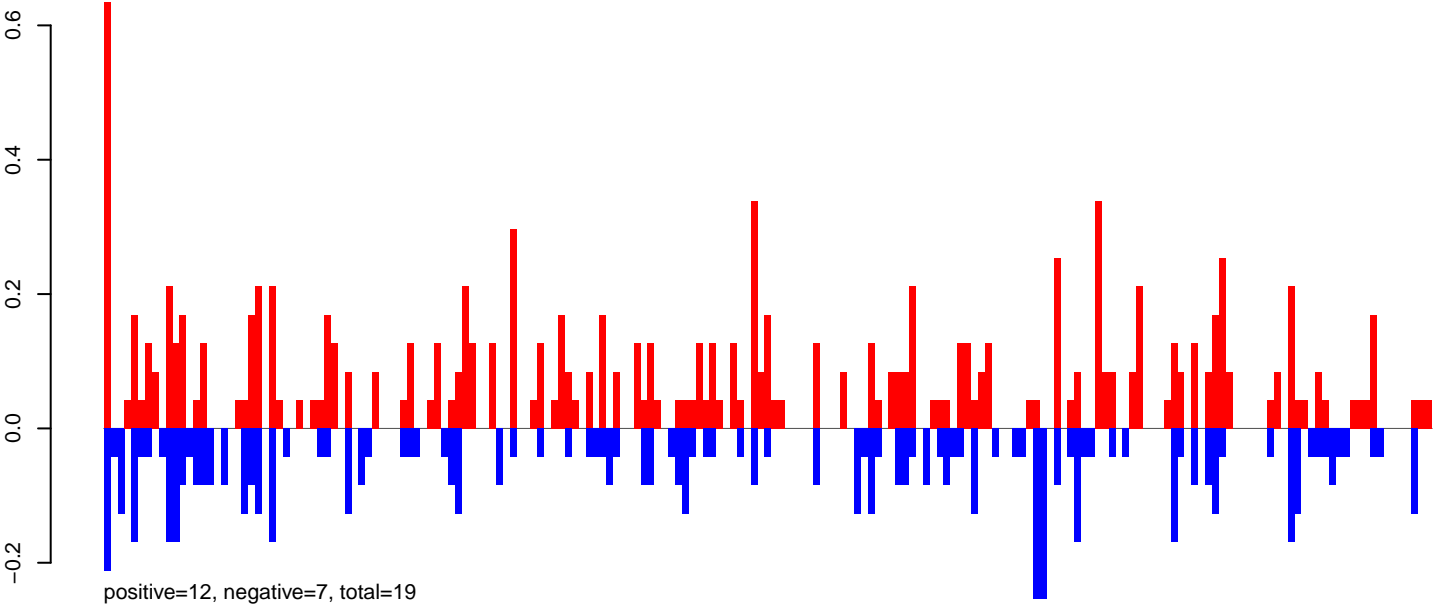
AeAlbo_C636_cells.rep



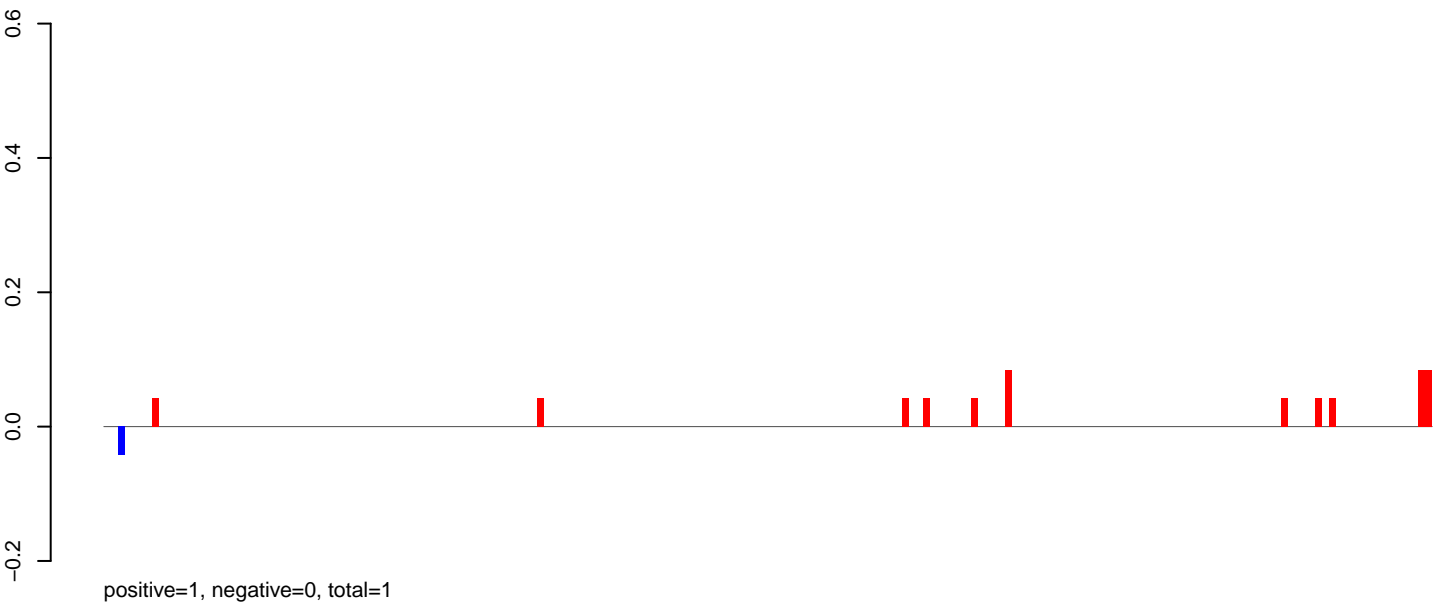
MSL000372-25.dna#478987.pl#6169
MH550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

0 1000 2000 3000 4000

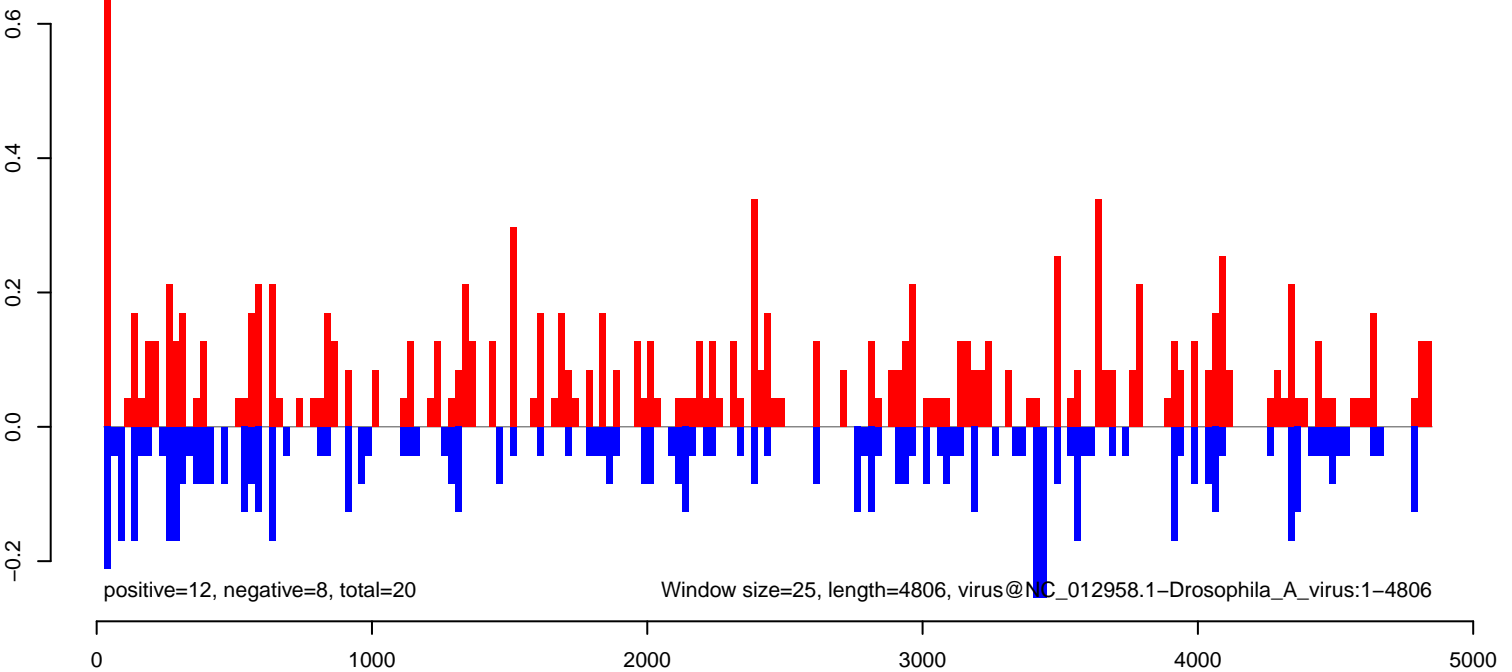
AeAlbo_C636_cells.18_23.rep



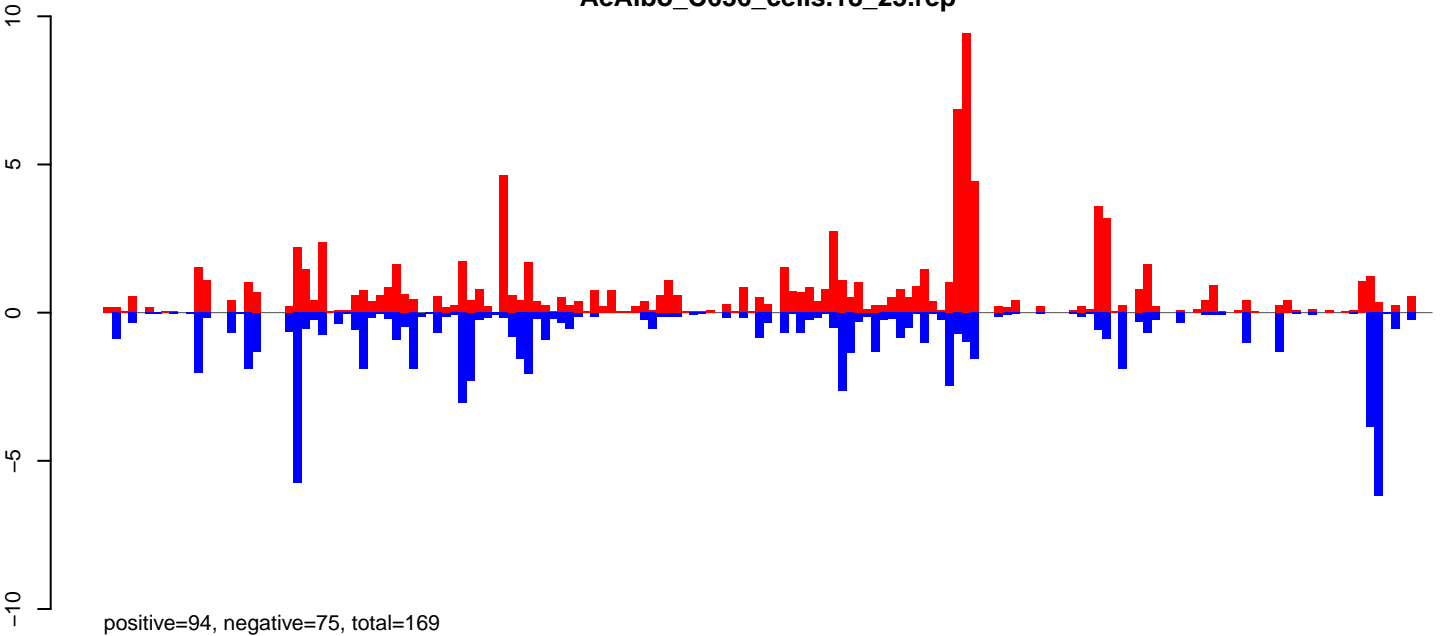
AeAlbo_C636_cells.24_35.rep



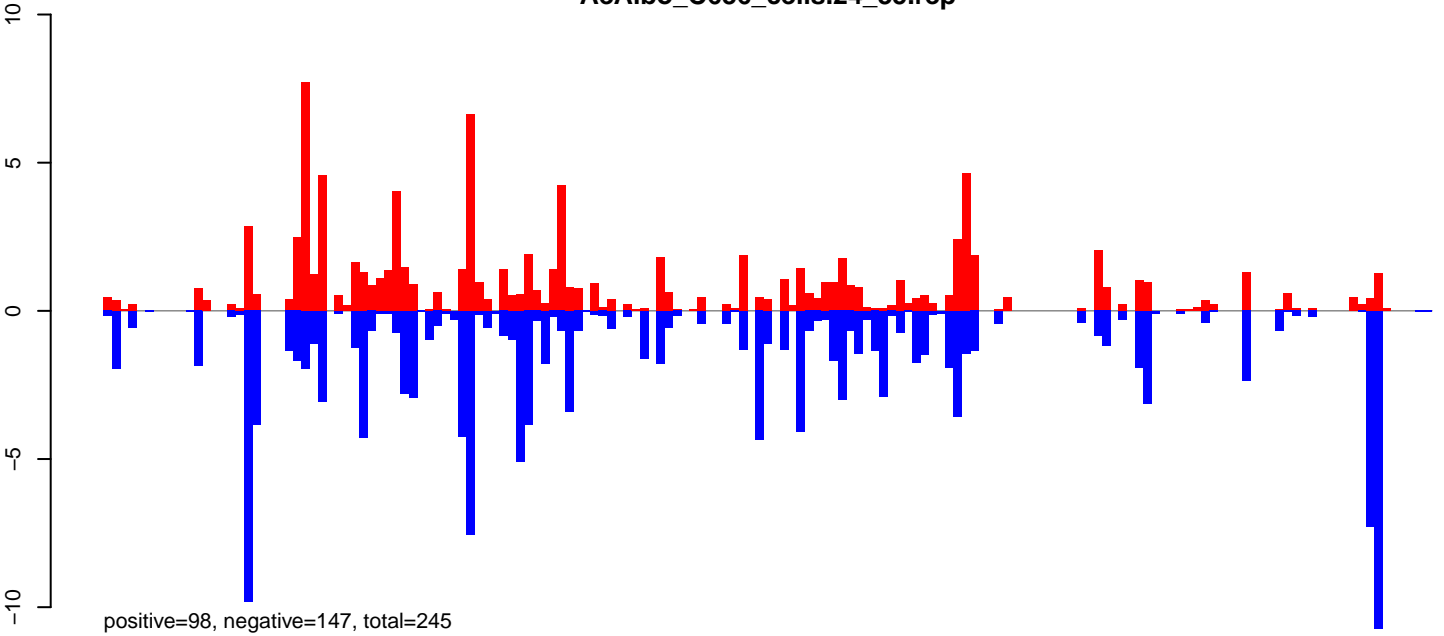
AeAlbo_C636_cells.rep



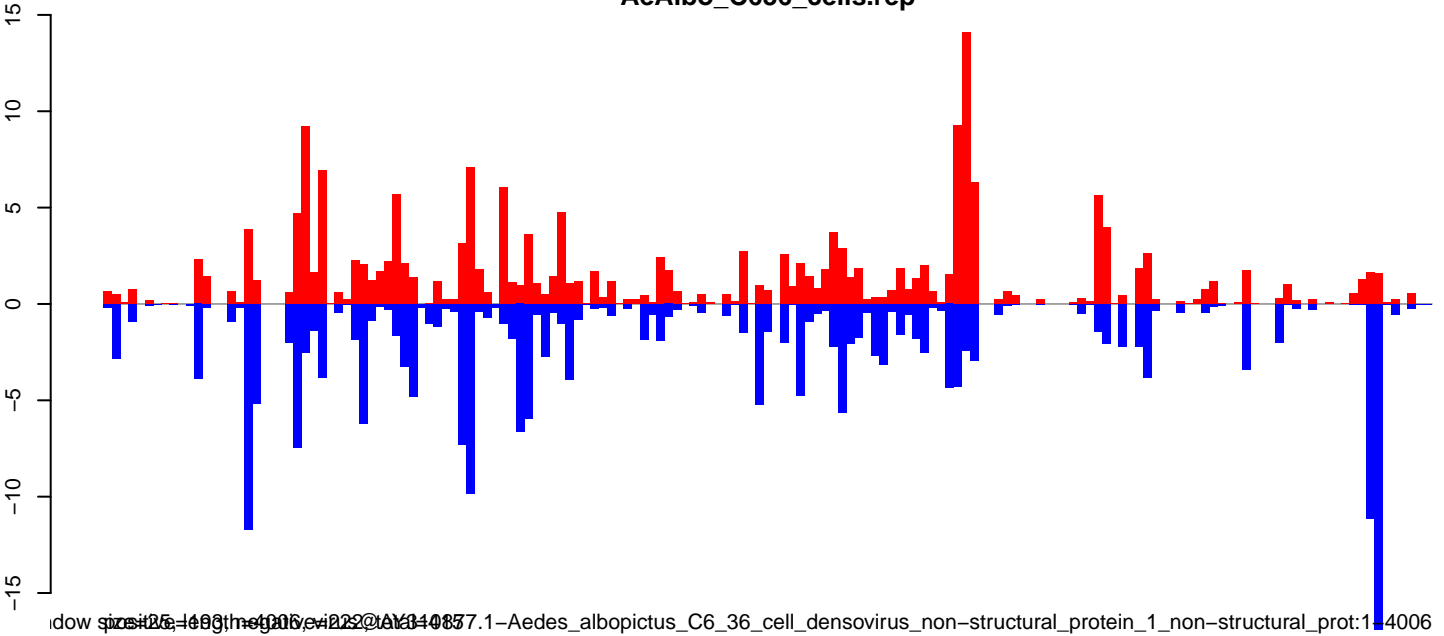
AeAlbo_C636_cells.18_23.rep



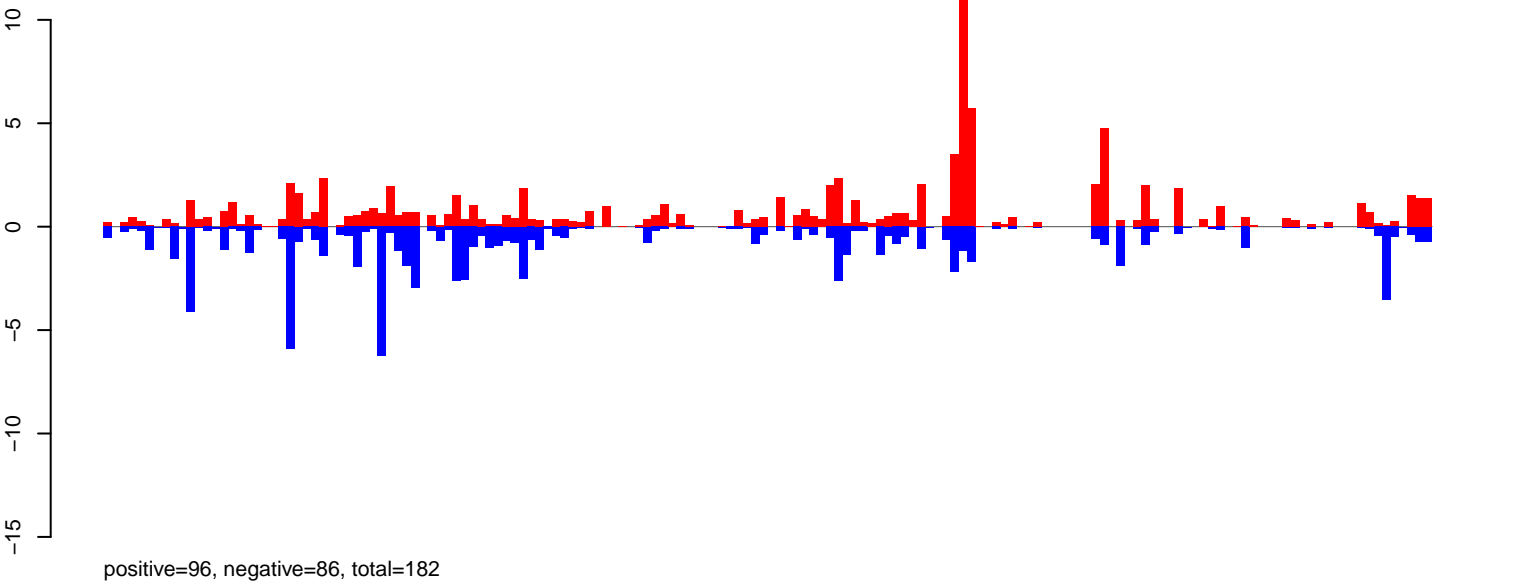
AeAlbo_C636_cells.24_35.rep



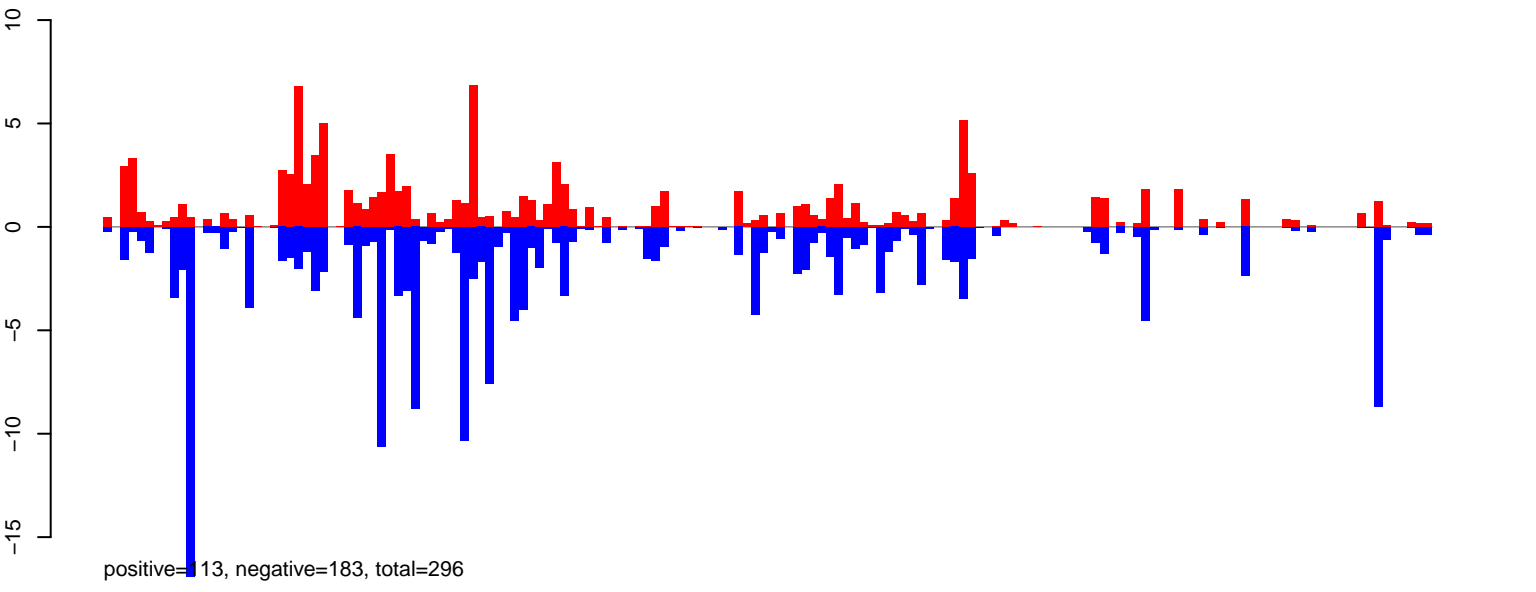
AeAlbo_C636_cells.rep



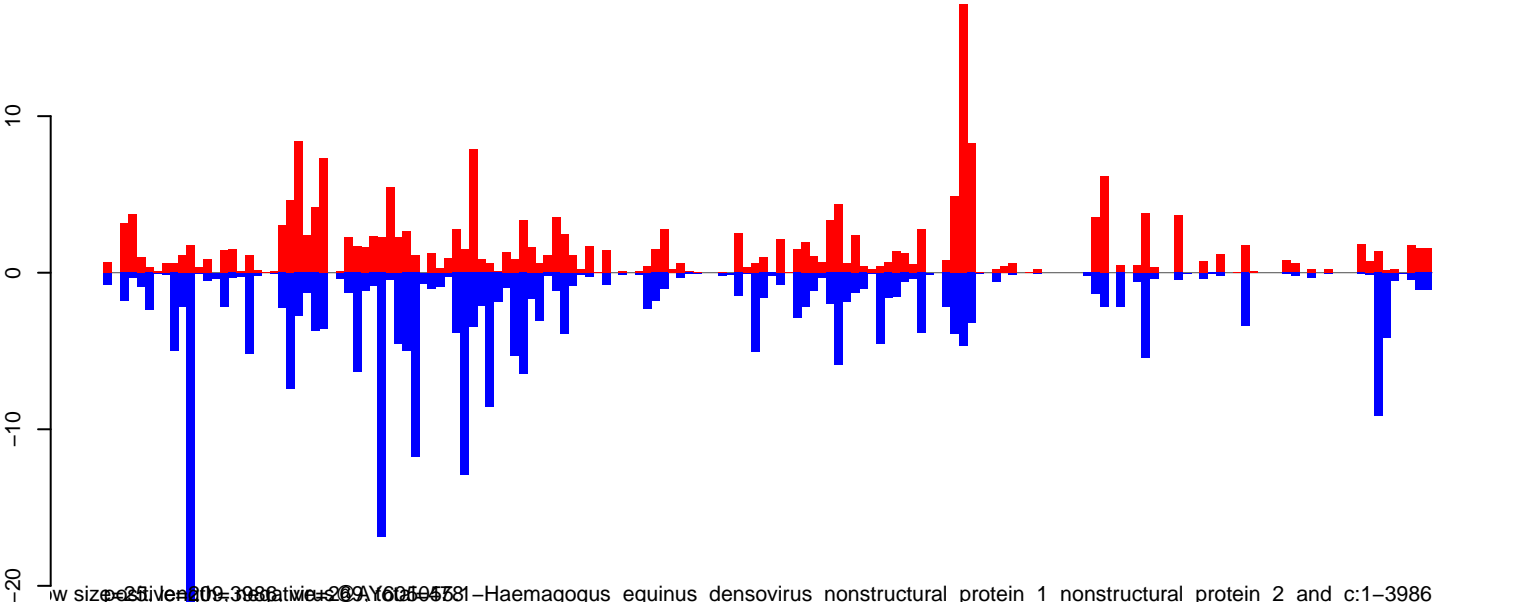
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



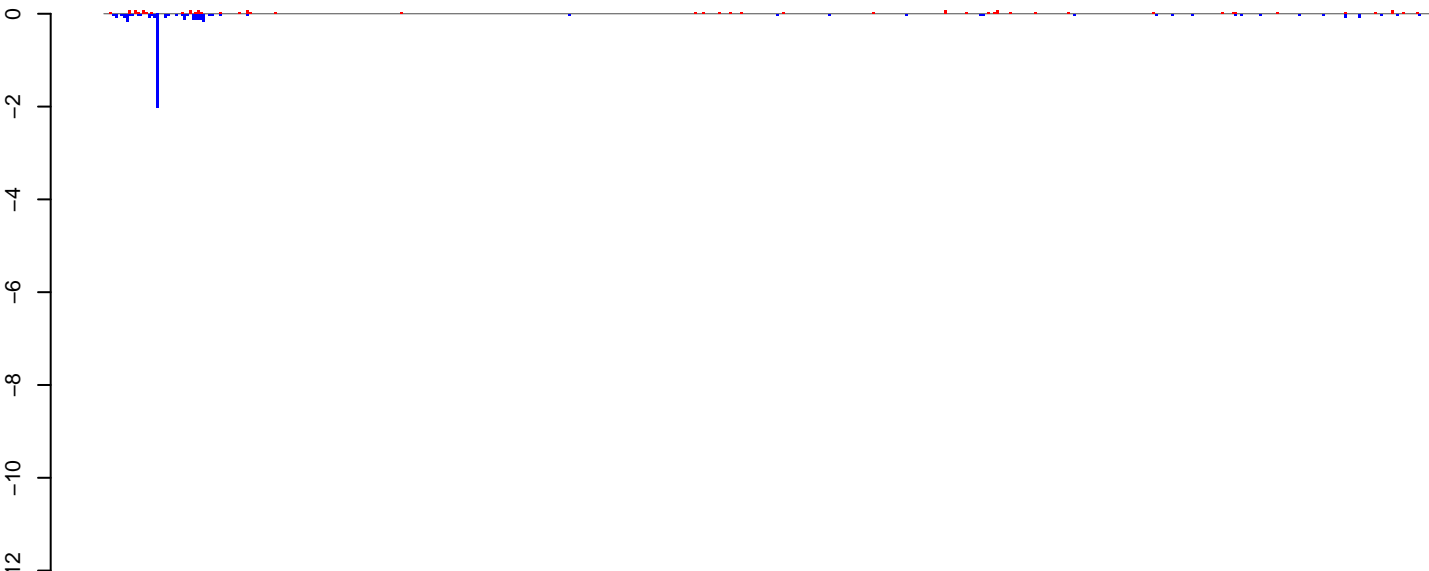
AeAlbo_C636_cells.rep



GenBank accession: AF029578.1 - Haemagogus equinus densovirus nonstructural protein 1 nonstructural protein 2 and c:1-3986

0 1000 2000 3000 4000

AeAlbo_C636_cells.18_23.rep



positive=2, negative=5, total=7

AeAlbo_C636_cells.24_35.rep



positive=2, negative=31, total=33

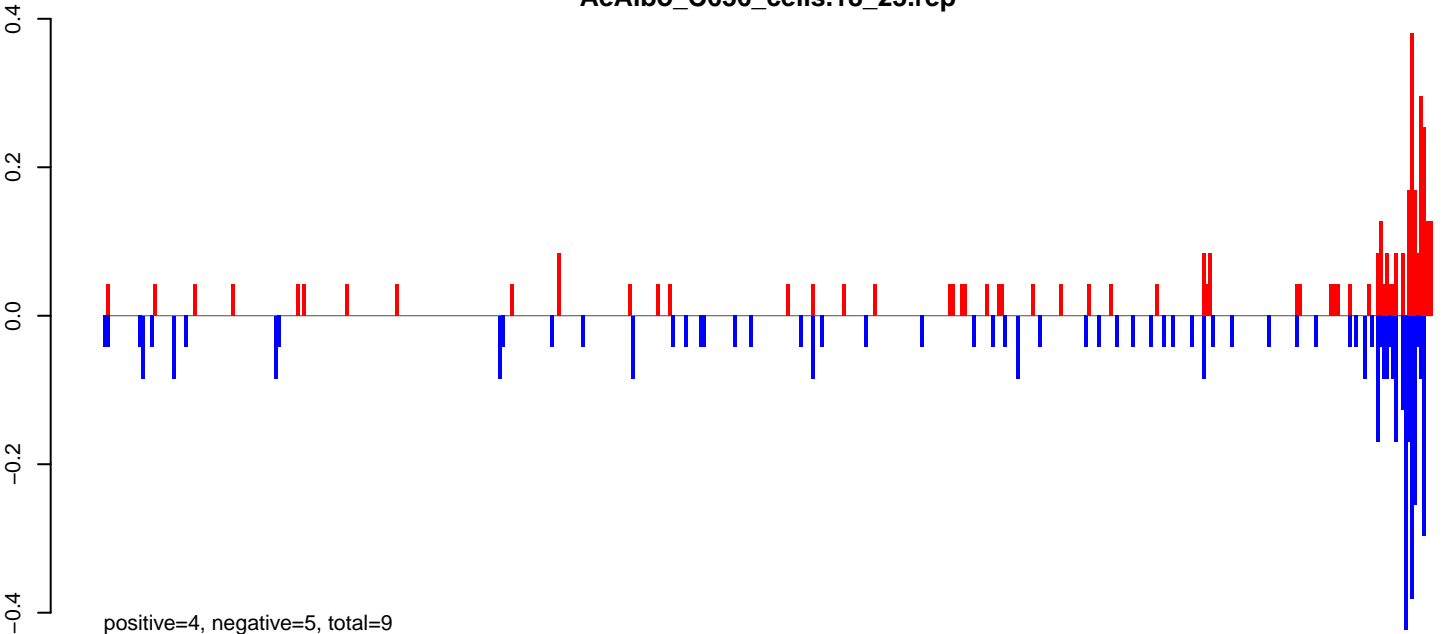
AeAlbo_C636_cells.rep



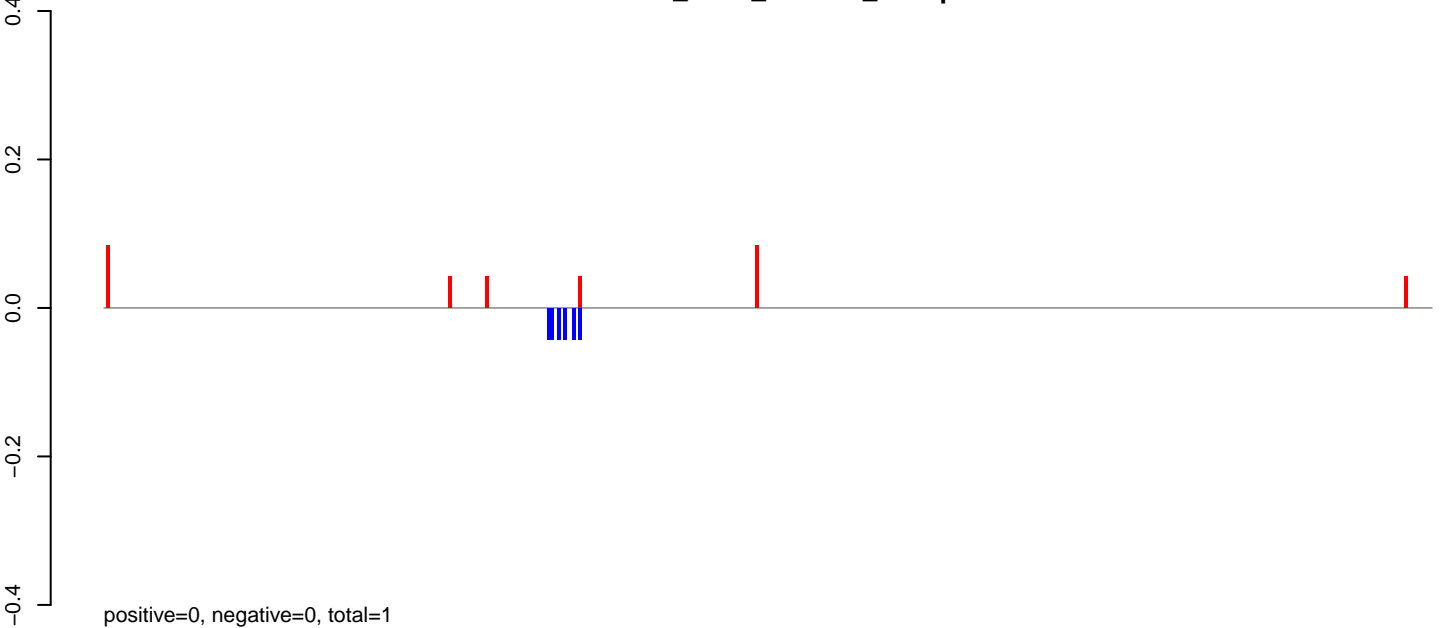
=25, length=12099, positive=25, negative=100, total=125
1-Phasi Charoen-like_virus_isolate_Rio_segment_S_plus_segment_M_plus_segment_L_Nucle:1-12099

0 2000 4000 6000 8000 10000 12000

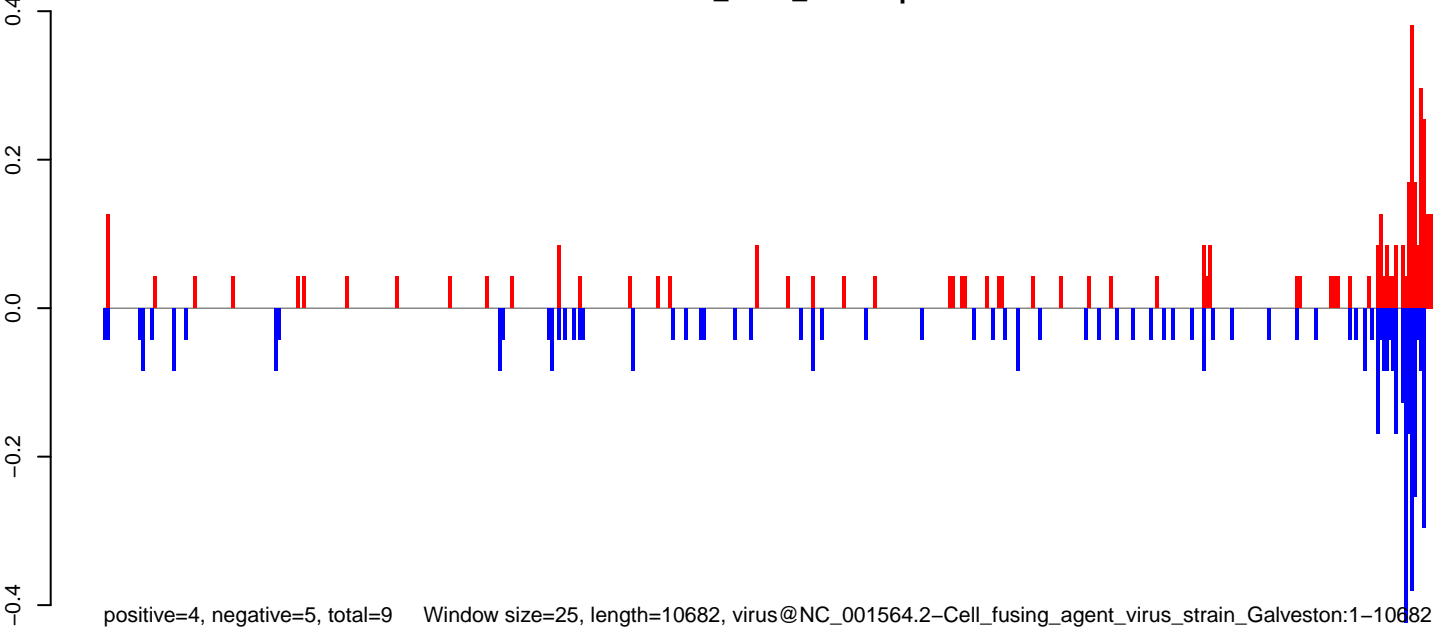
AeAlbo_C636_cells.18_23.rep



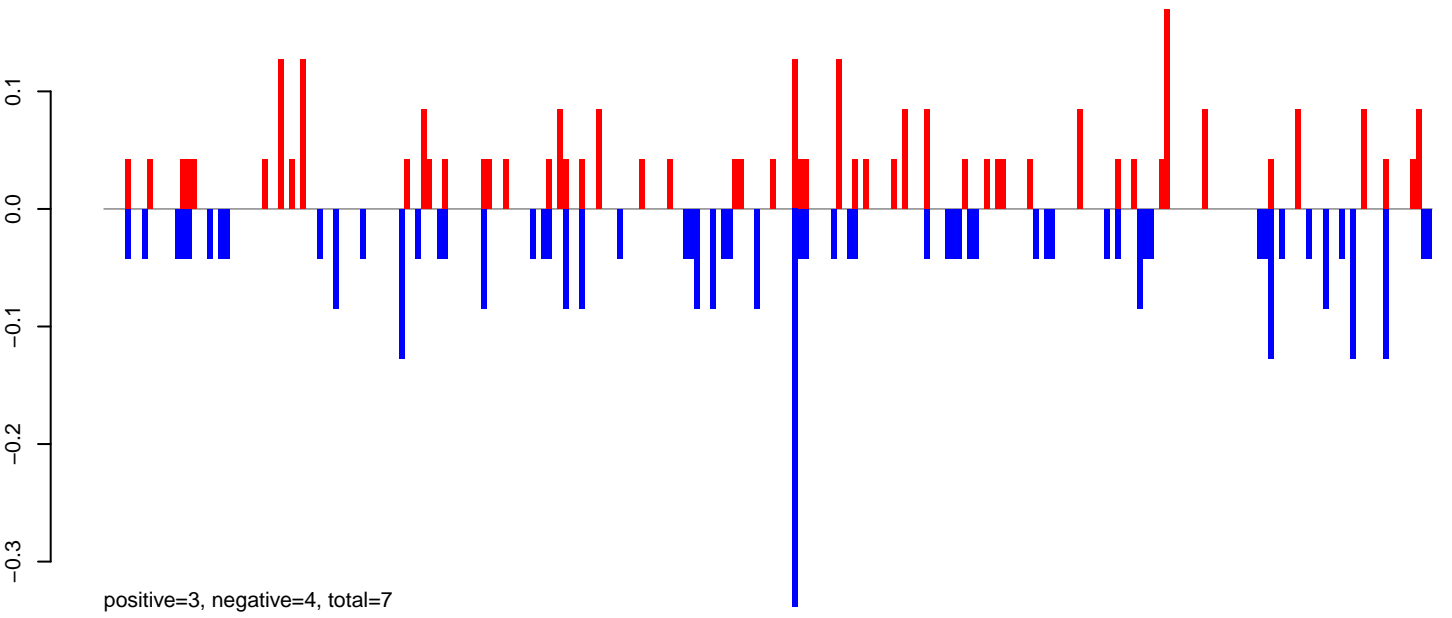
AeAlbo_C636_cells.24_35.rep



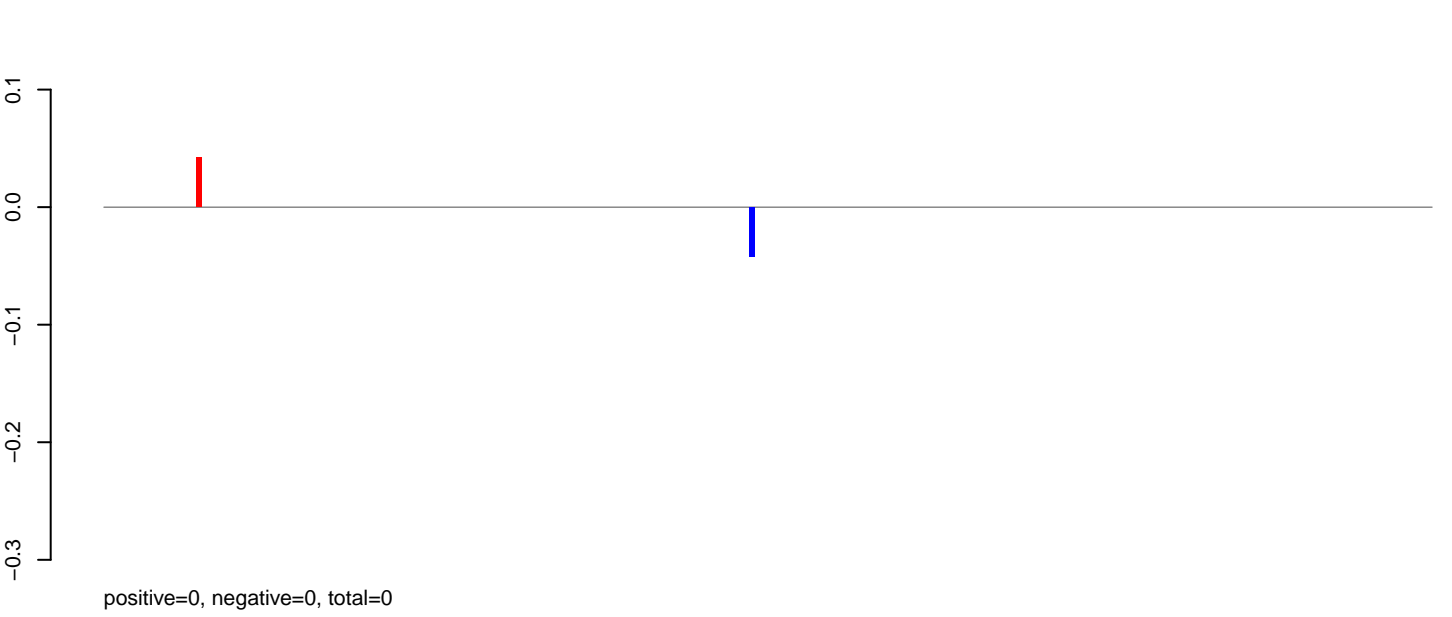
AeAlbo_C636_cells.rep



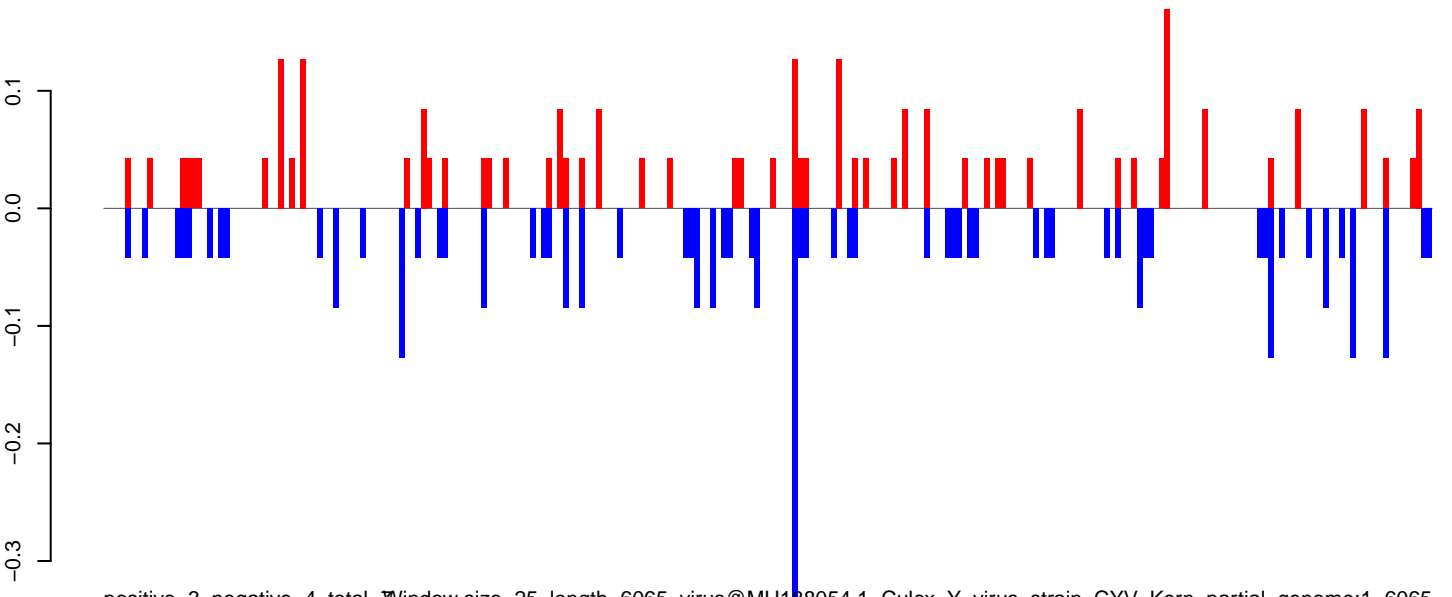
AeAlbo_C636_cells.18_23.rep



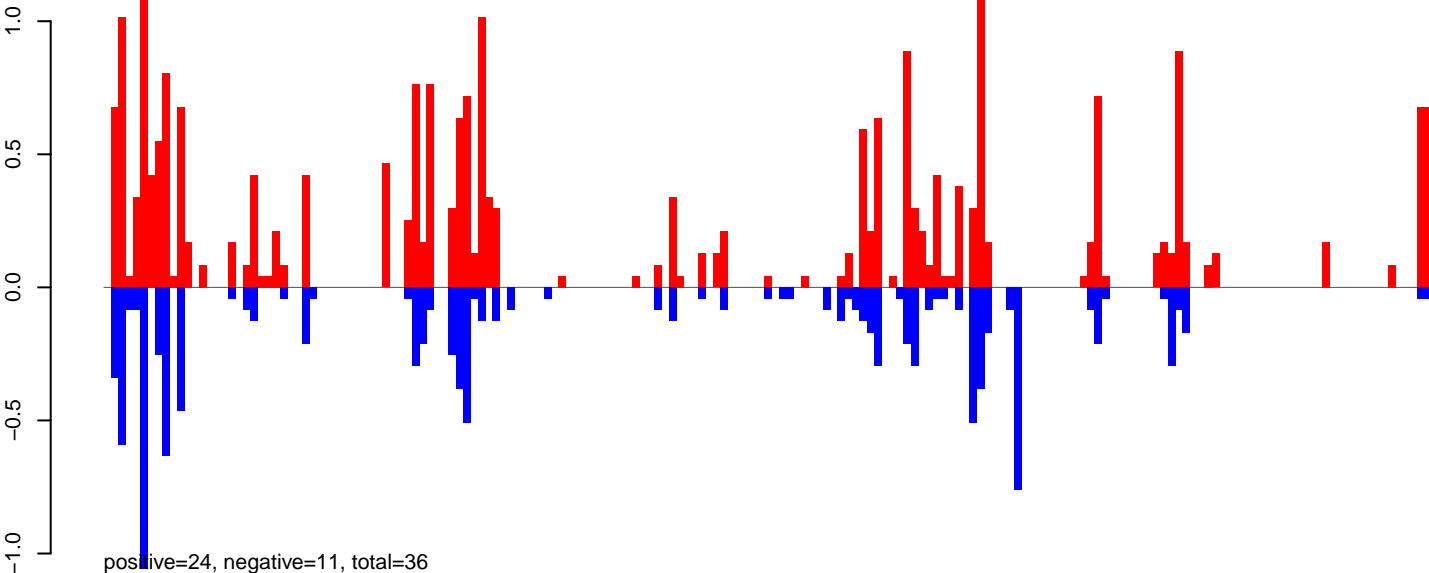
AeAlbo_C636_cells.24_35.rep



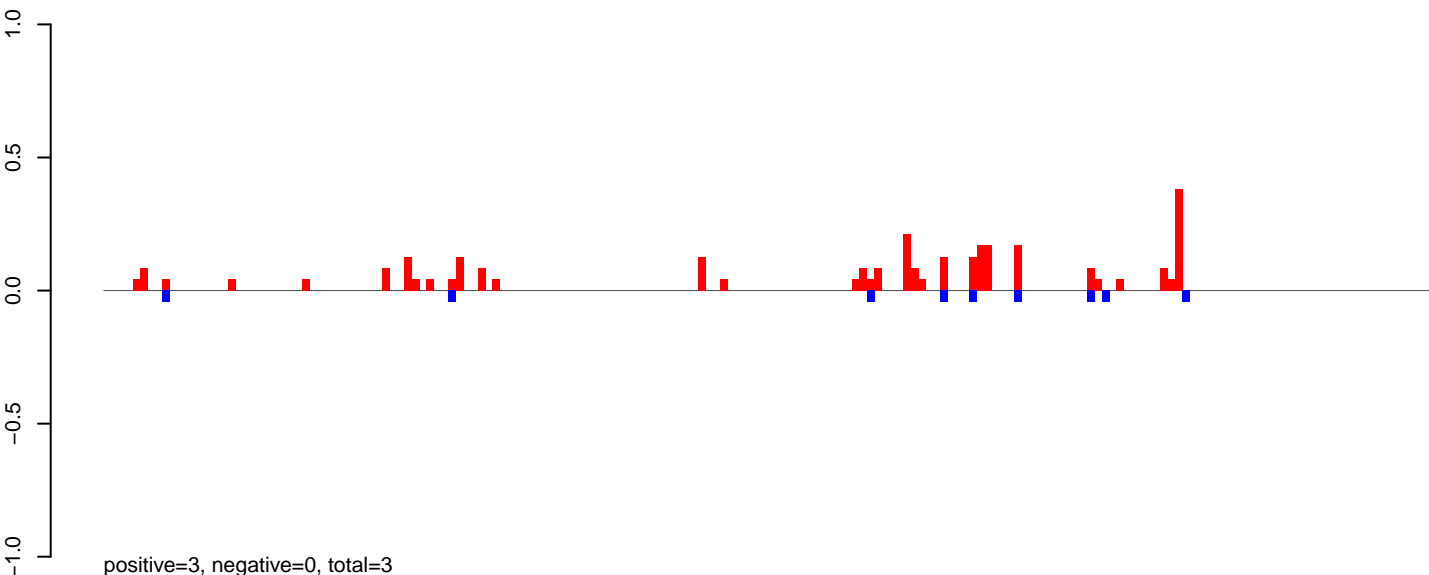
AeAlbo_C636_cells.rep



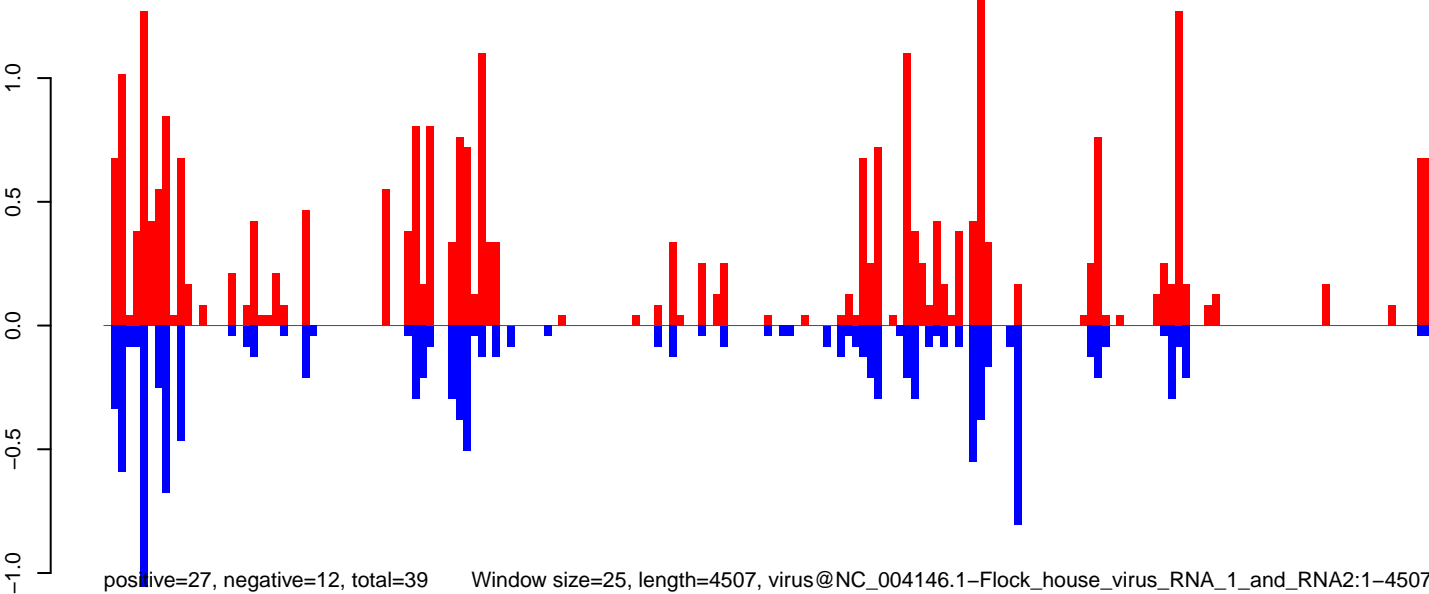
AeAlbo_C636_cells.18_23.rep



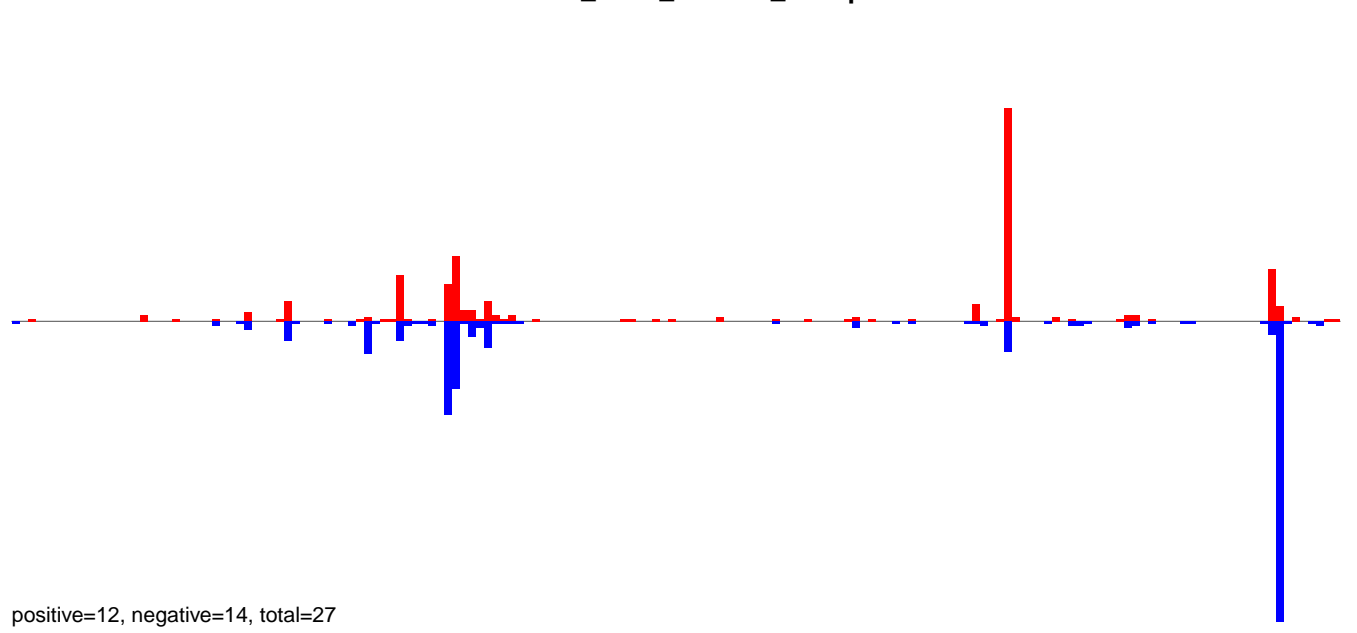
AeAlbo_C636_cells.24_35.rep



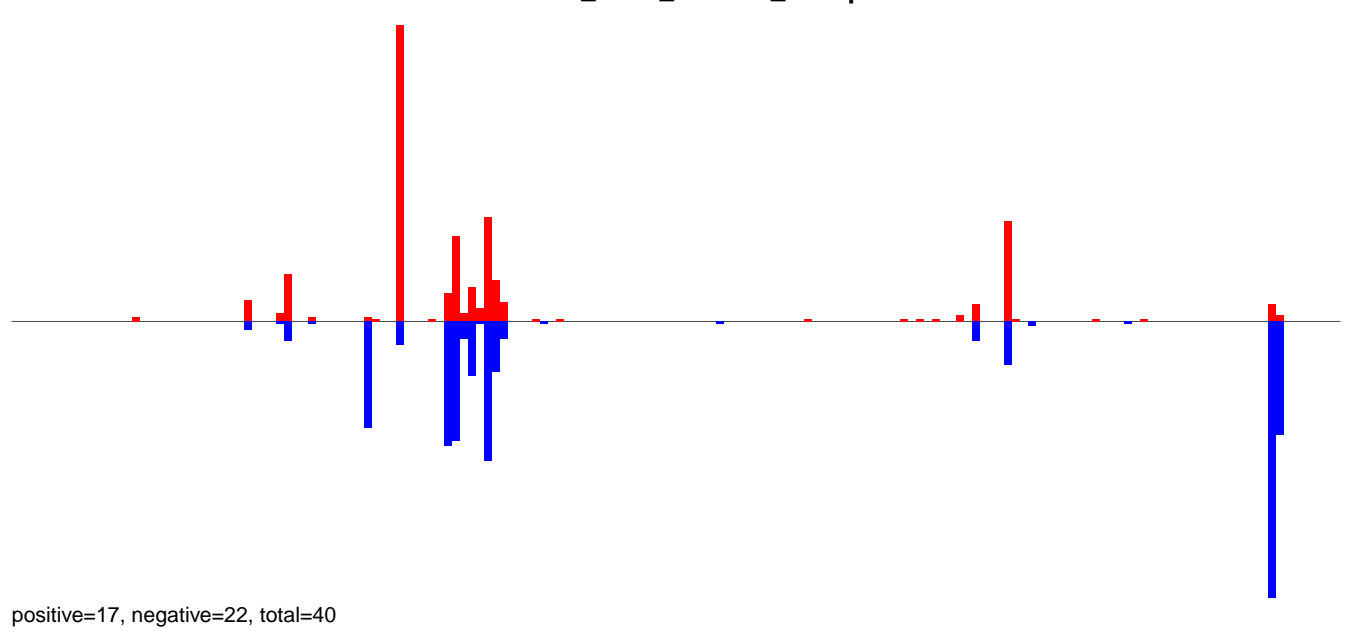
AeAlbo_C636_cells.rep



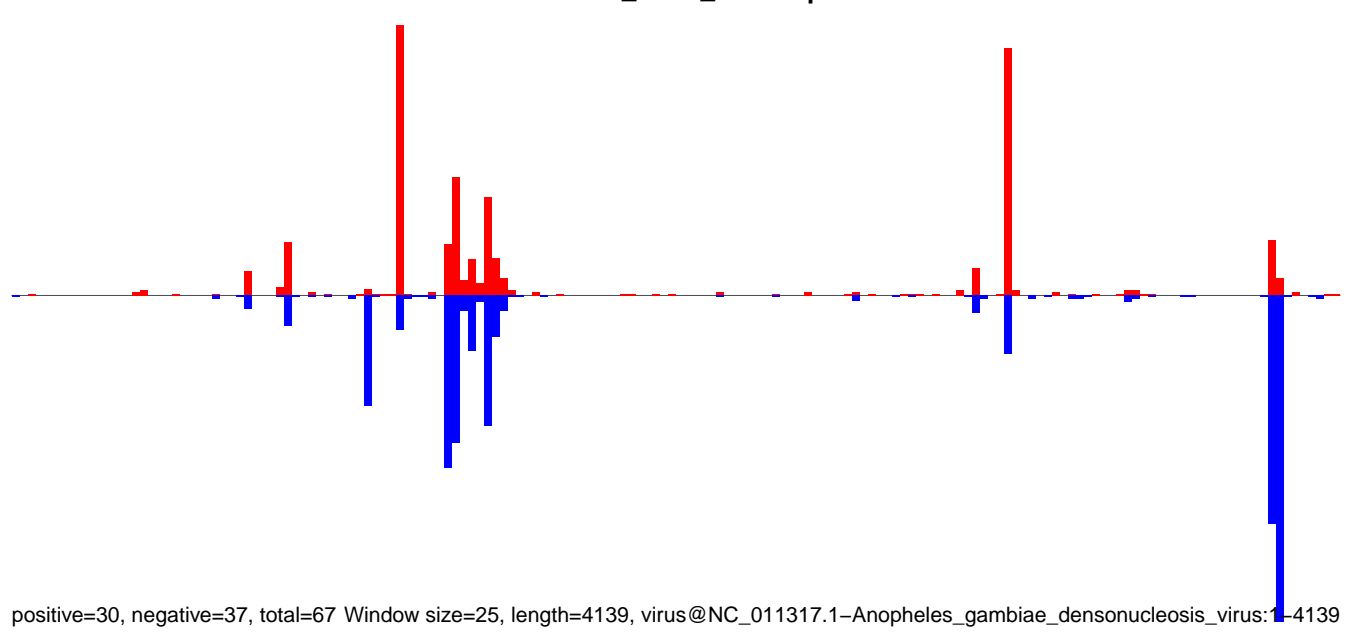
AeAlbo_C636_cells.18_23.rep



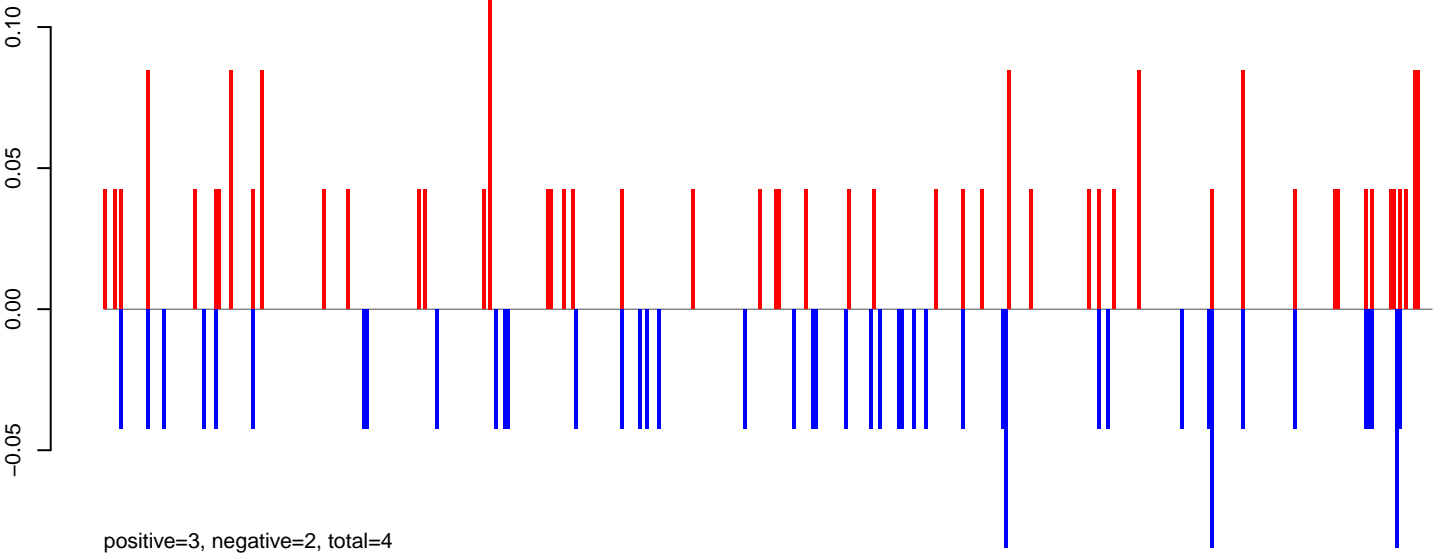
AeAlbo_C636_cells.24_35.rep



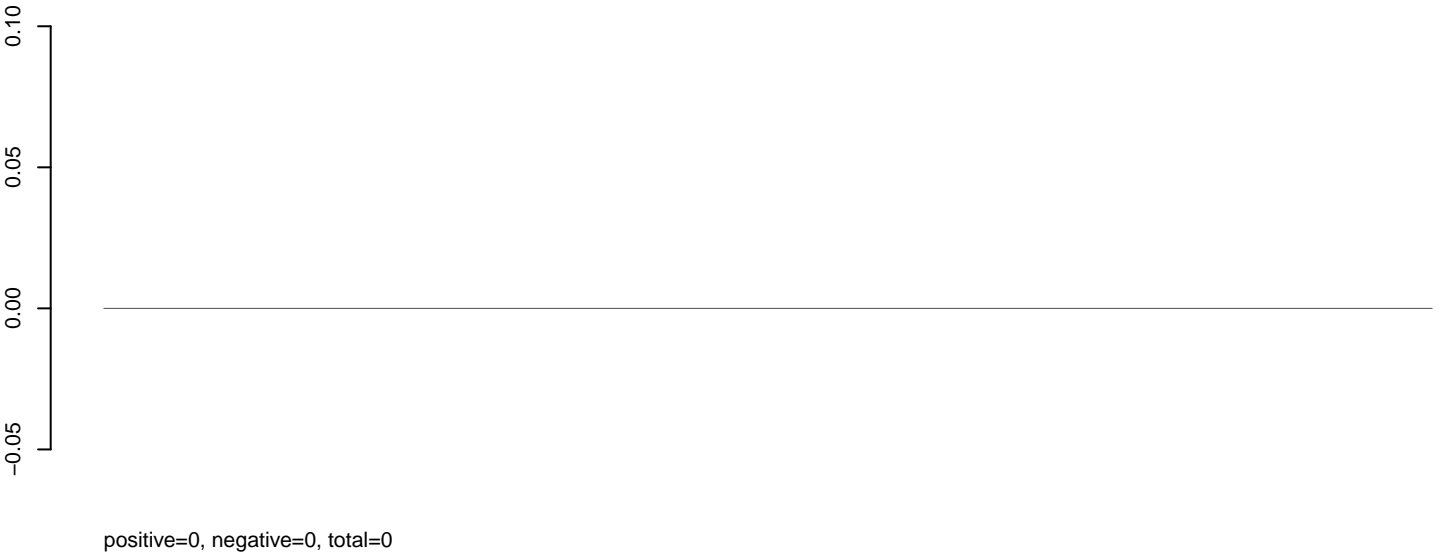
AeAlbo_C636_cells.rep



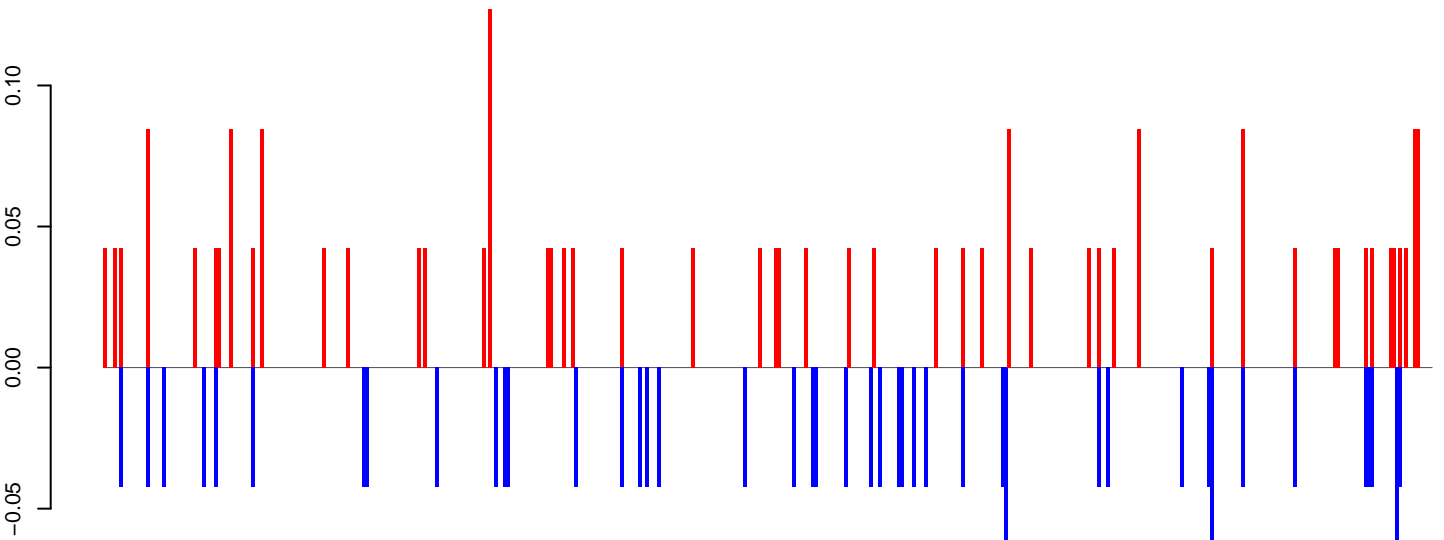
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



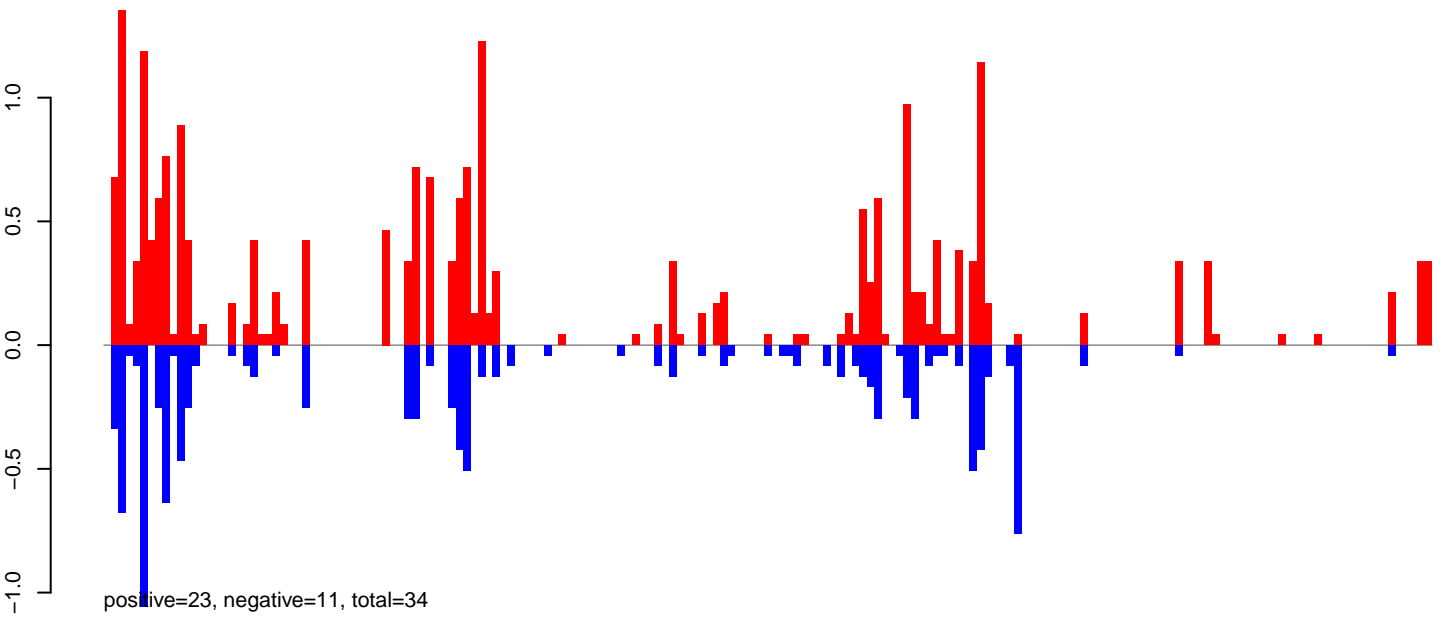
AeAlbo_C636_cells.rep



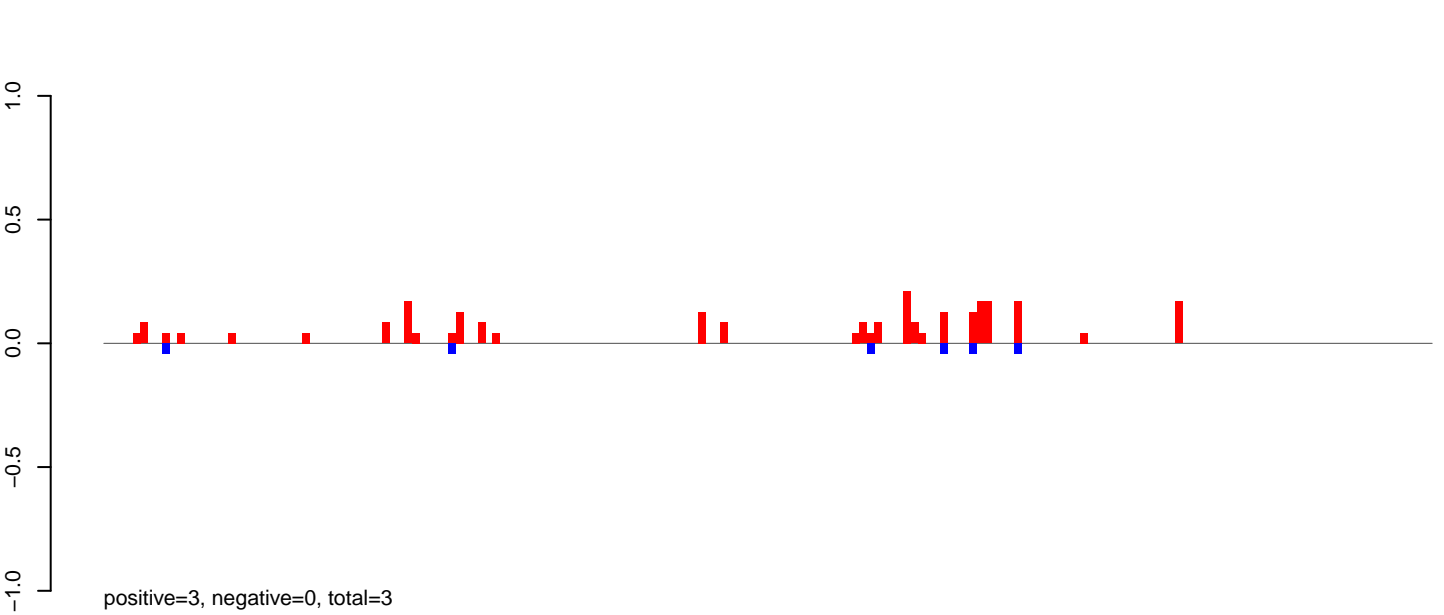
Window size=25, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

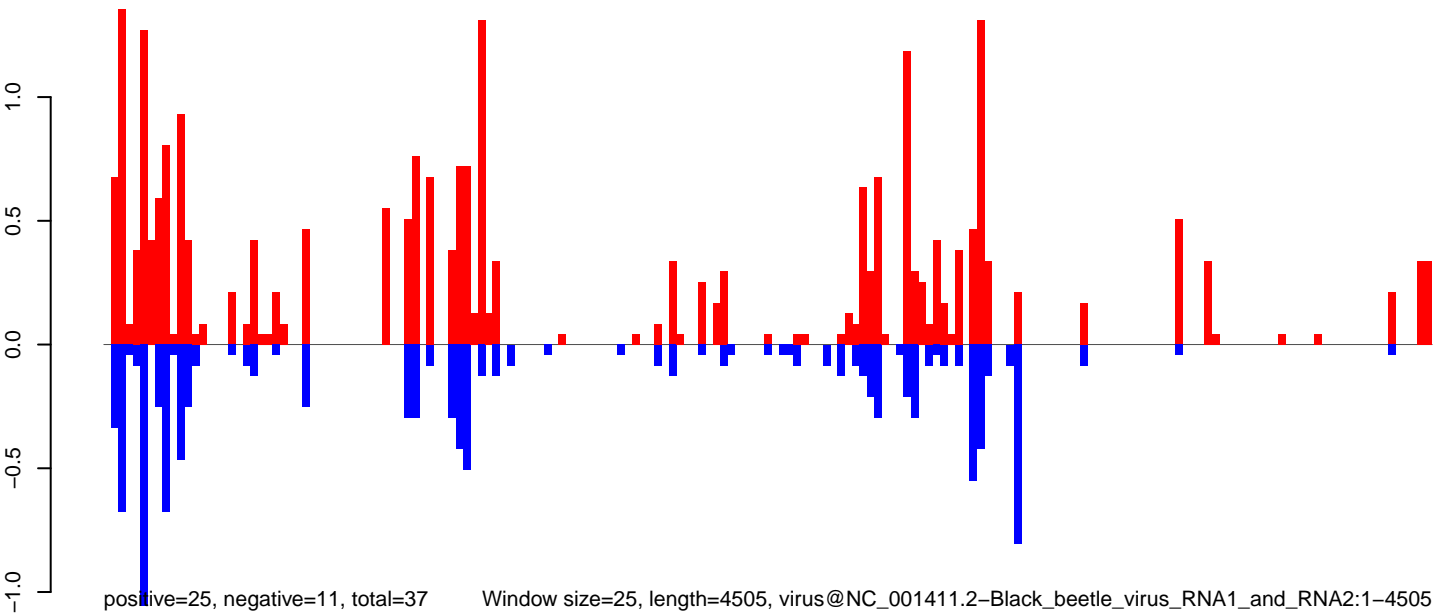
AeAlbo_C636_cells.18_23.rep



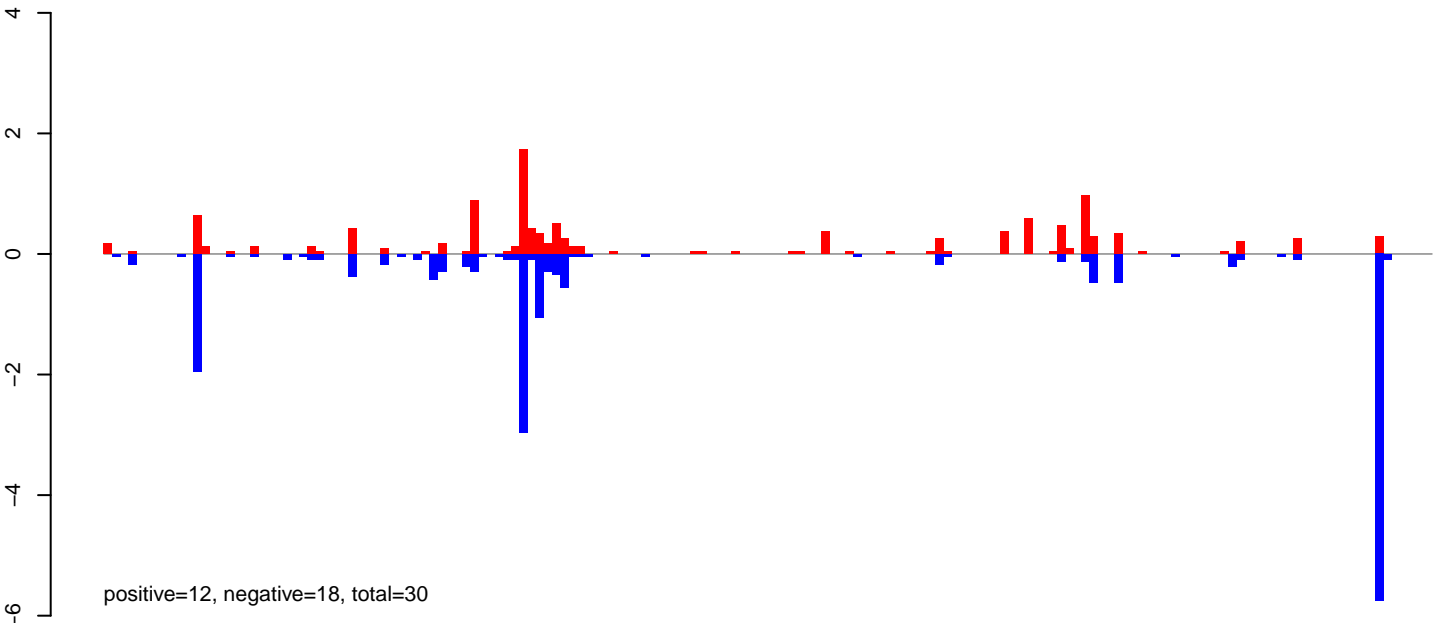
AeAlbo_C636_cells.24_35.rep



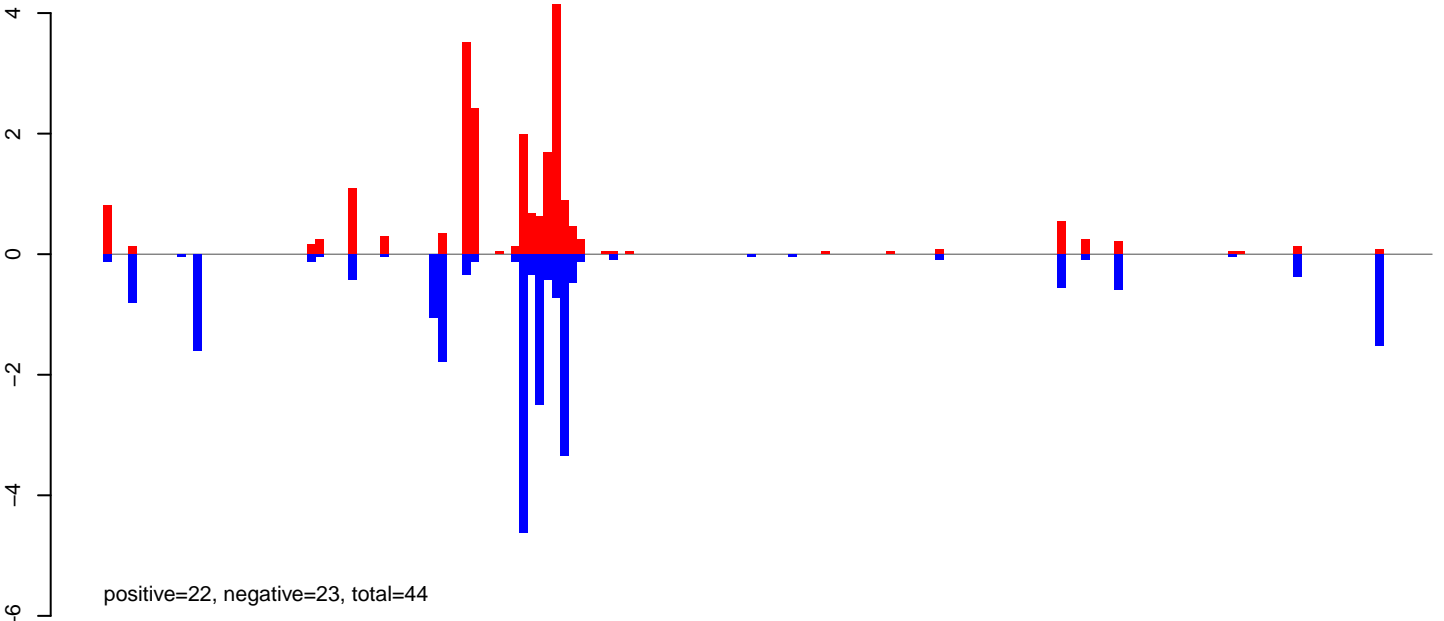
AeAlbo_C636_cells.rep



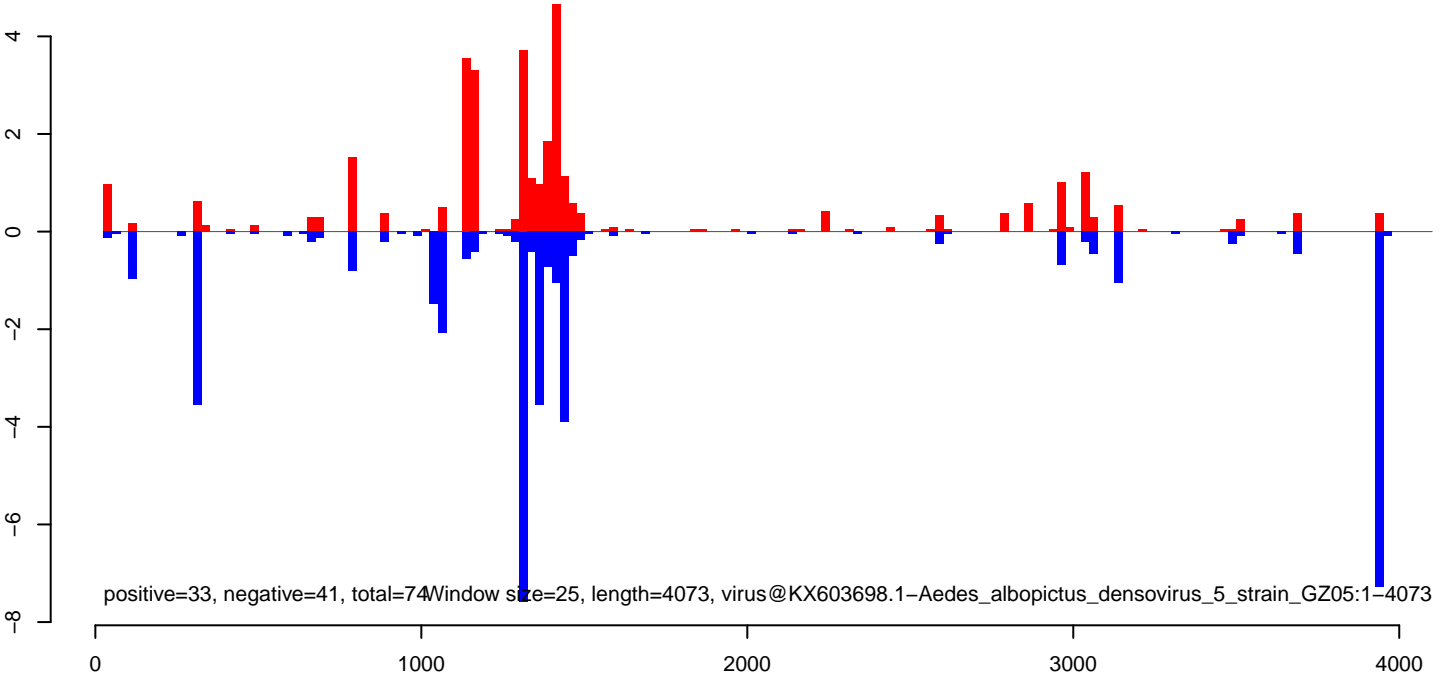
AeAlbo_C636_cells.18_23.rep



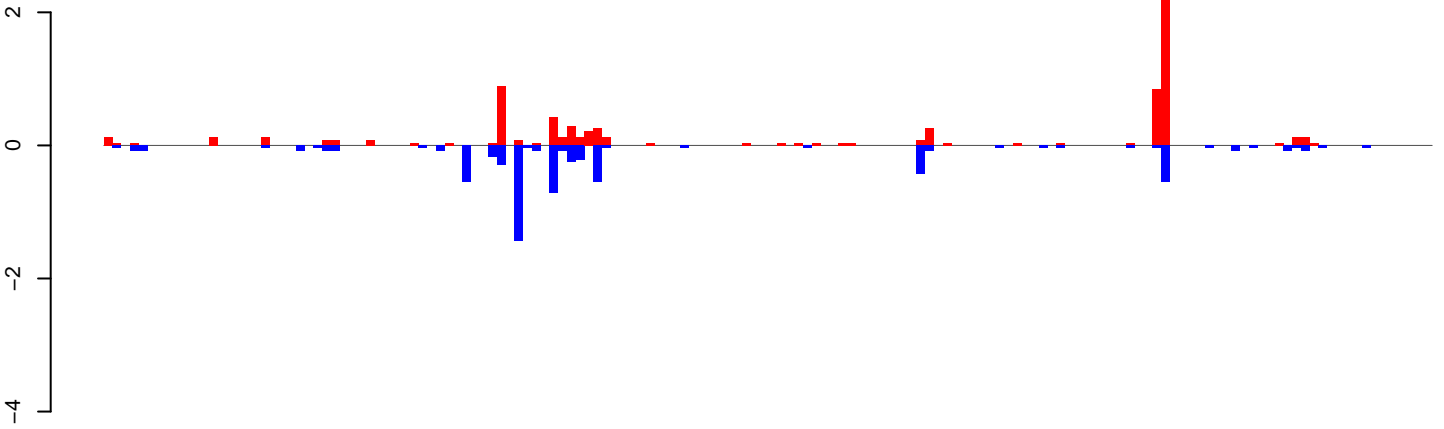
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

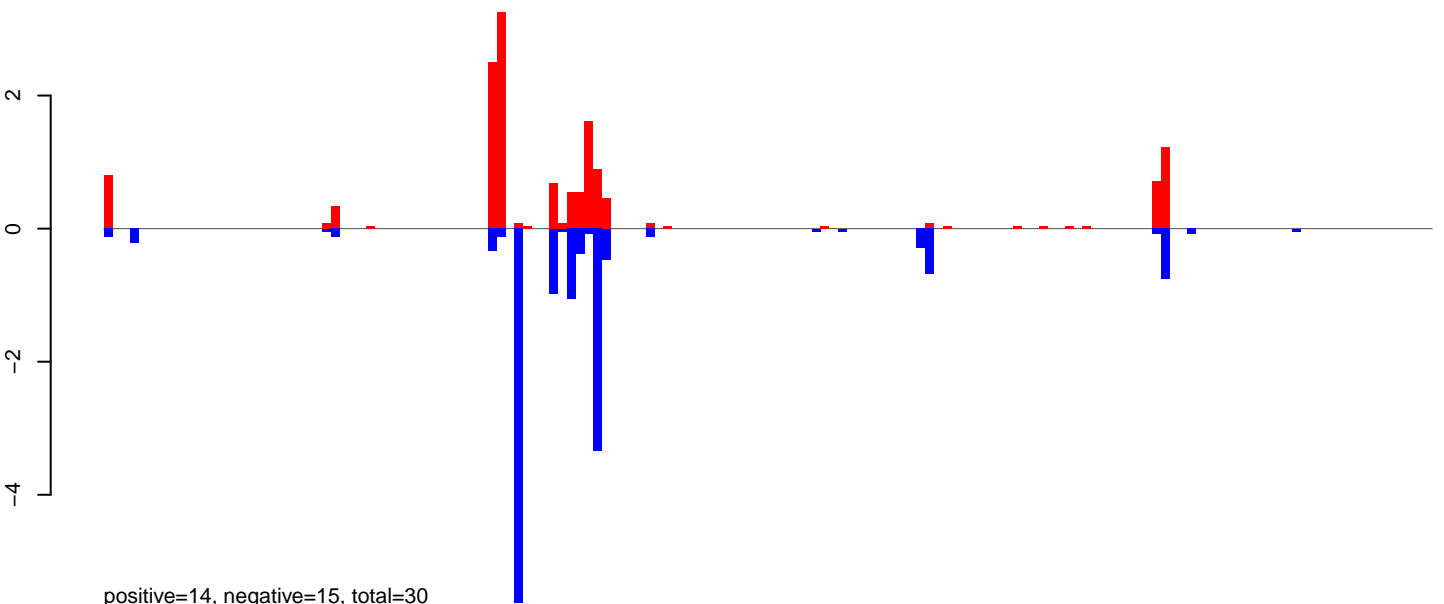


AeAlbo_C636_cells.18_23.rep



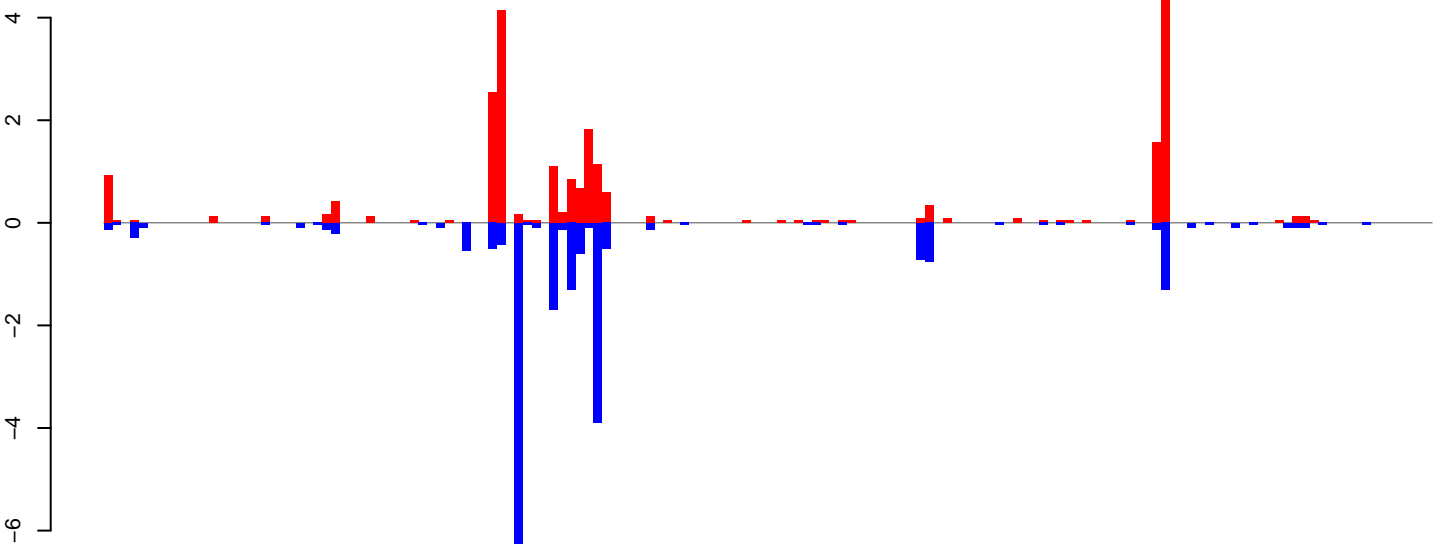
positive=9, negative=7, total=16

AeAlbo_C636_cells.24_35.rep



positive=14, negative=15, total=30

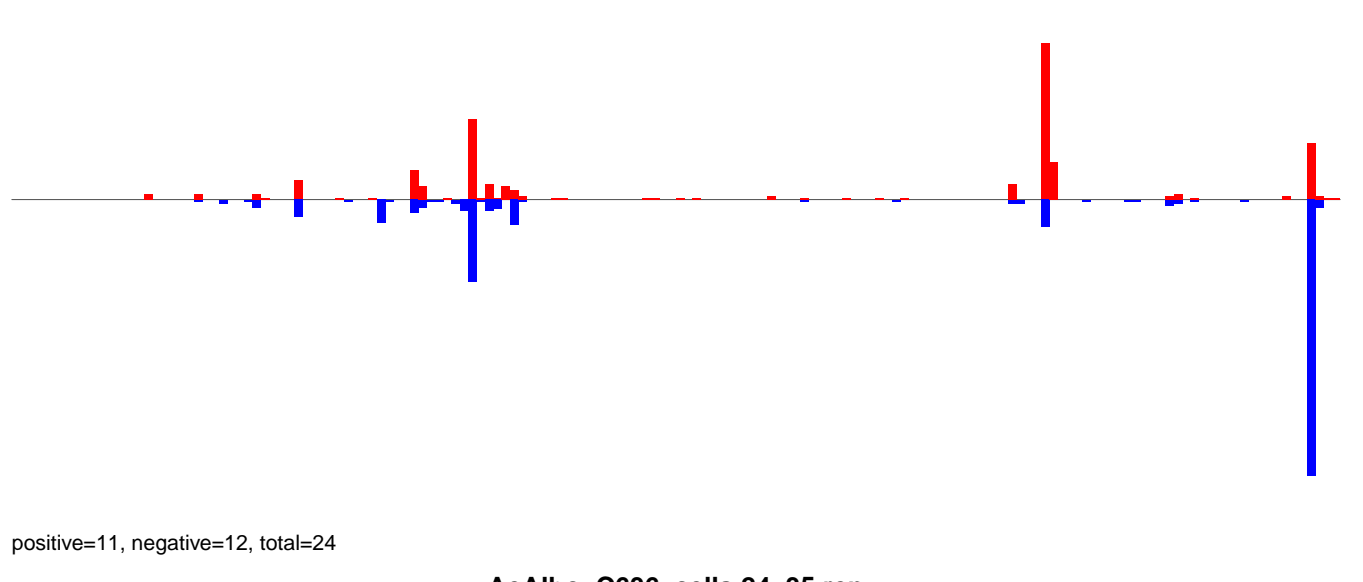
AeAlbo_C636_cells.rep



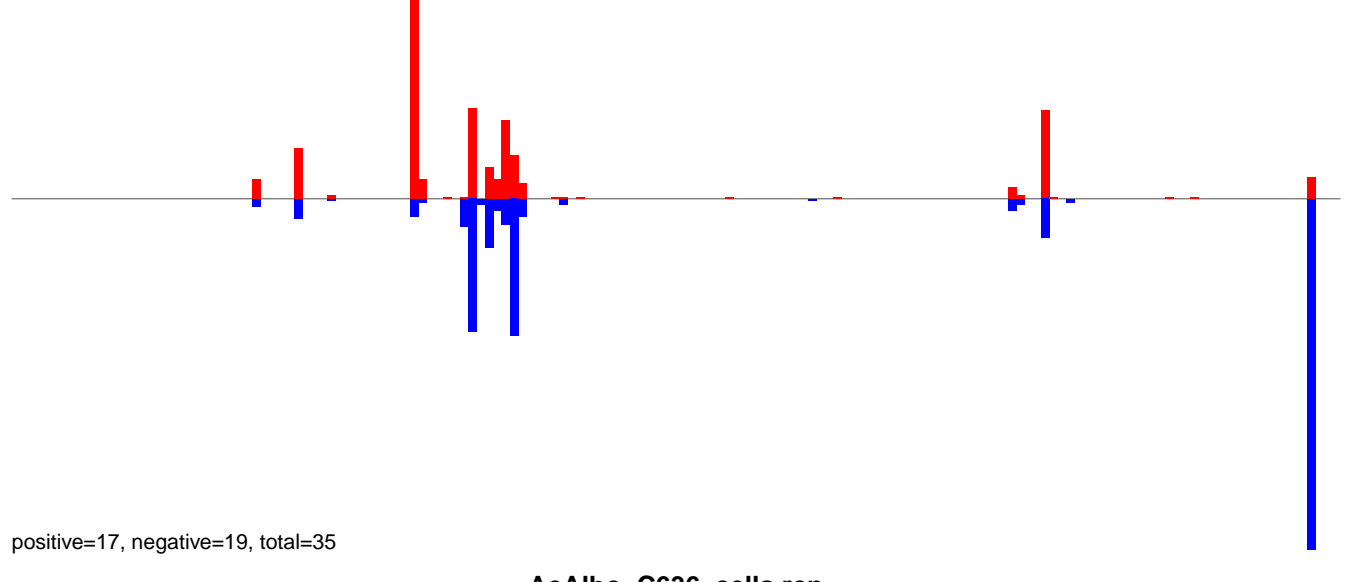
positive=23, negative=22, total=45 window size=25, length=3776, virus@NC_012636.1-Aedes_aegypti_densovirus_2_strain_0814616:1-3776

0 1000 2000 3000

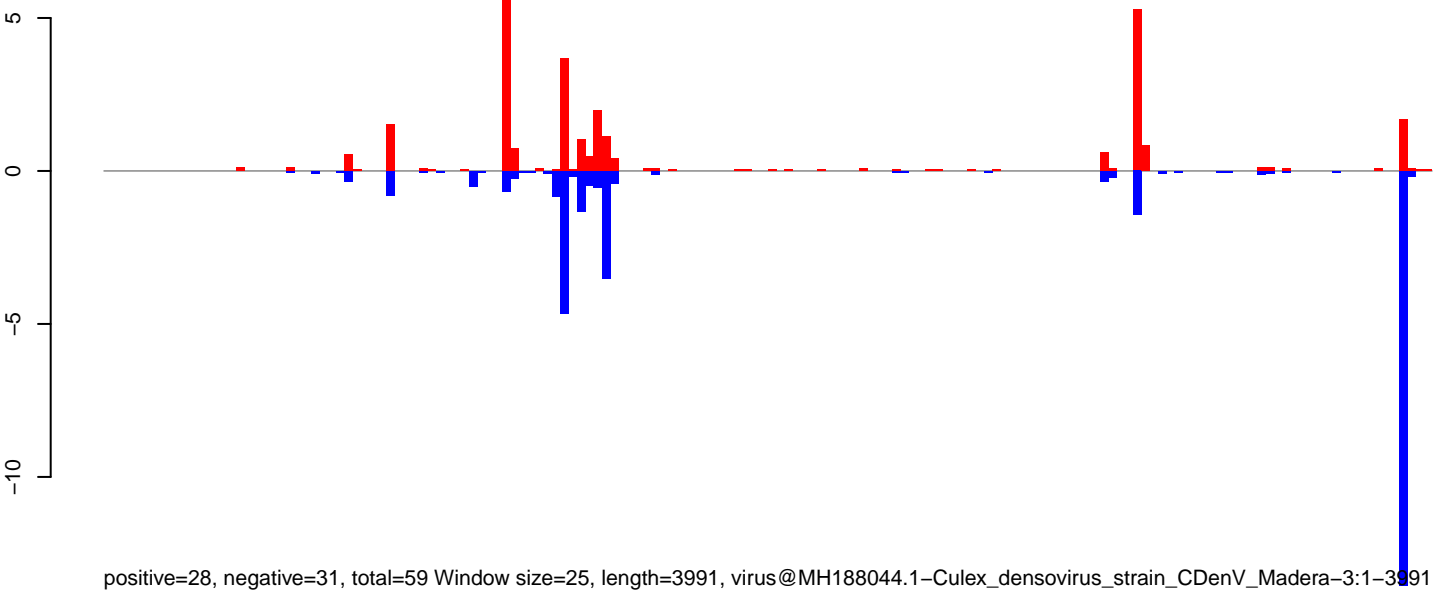
AeAlbo_C636_cells.18_23.rep



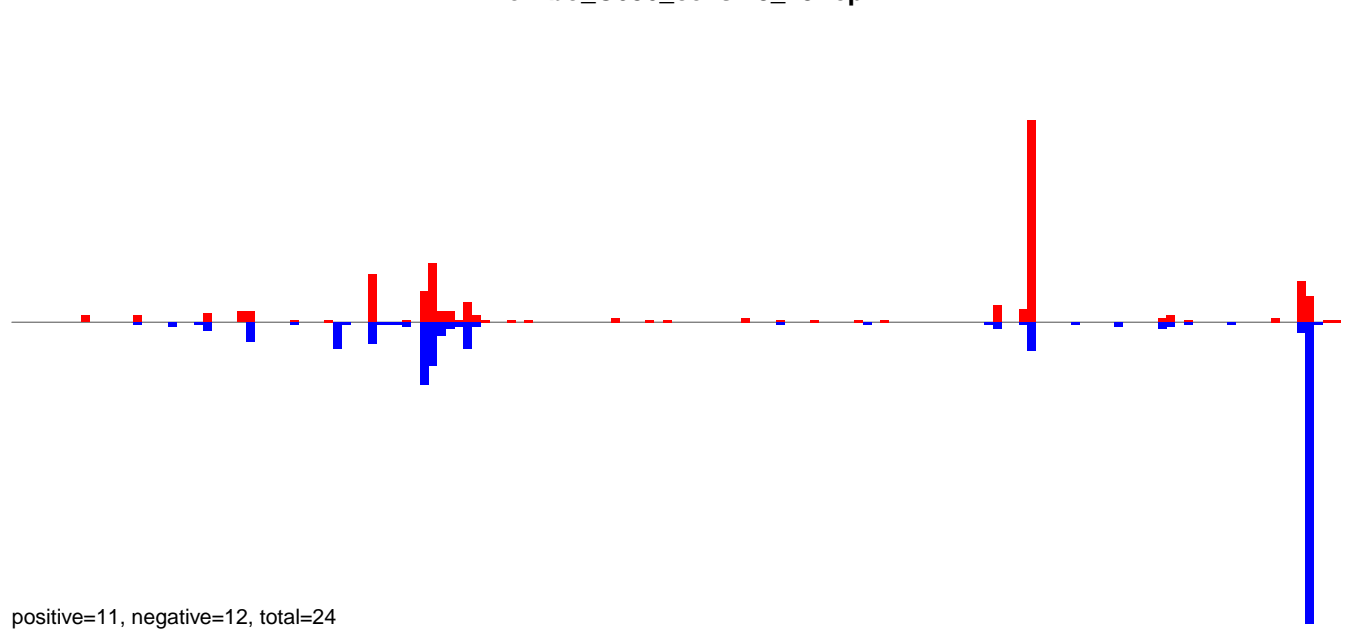
AeAlbo_C636_cells.24_35.rep



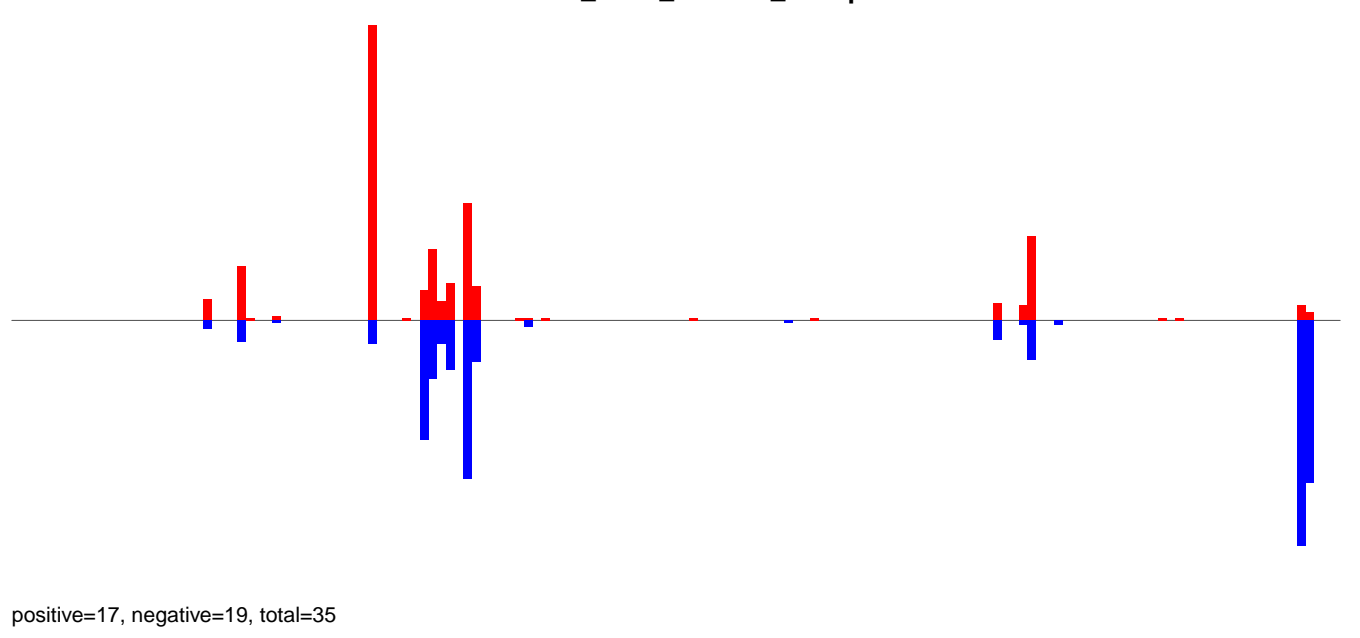
AeAlbo_C636_cells.rep



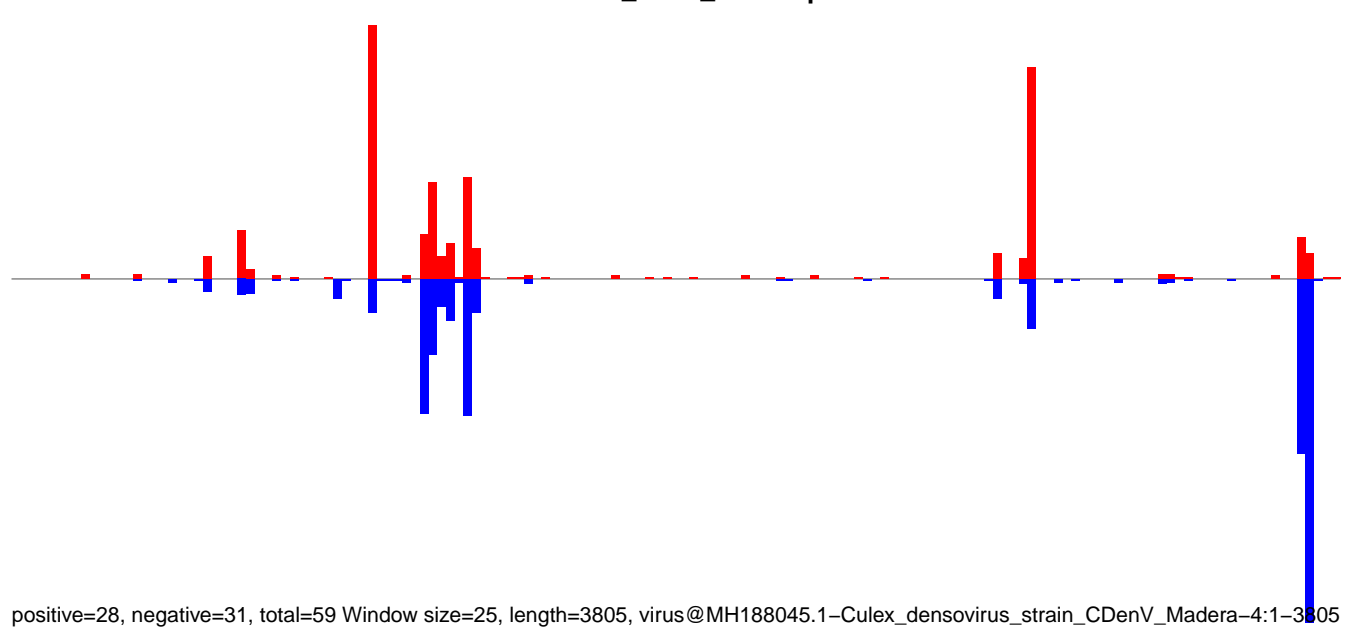
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

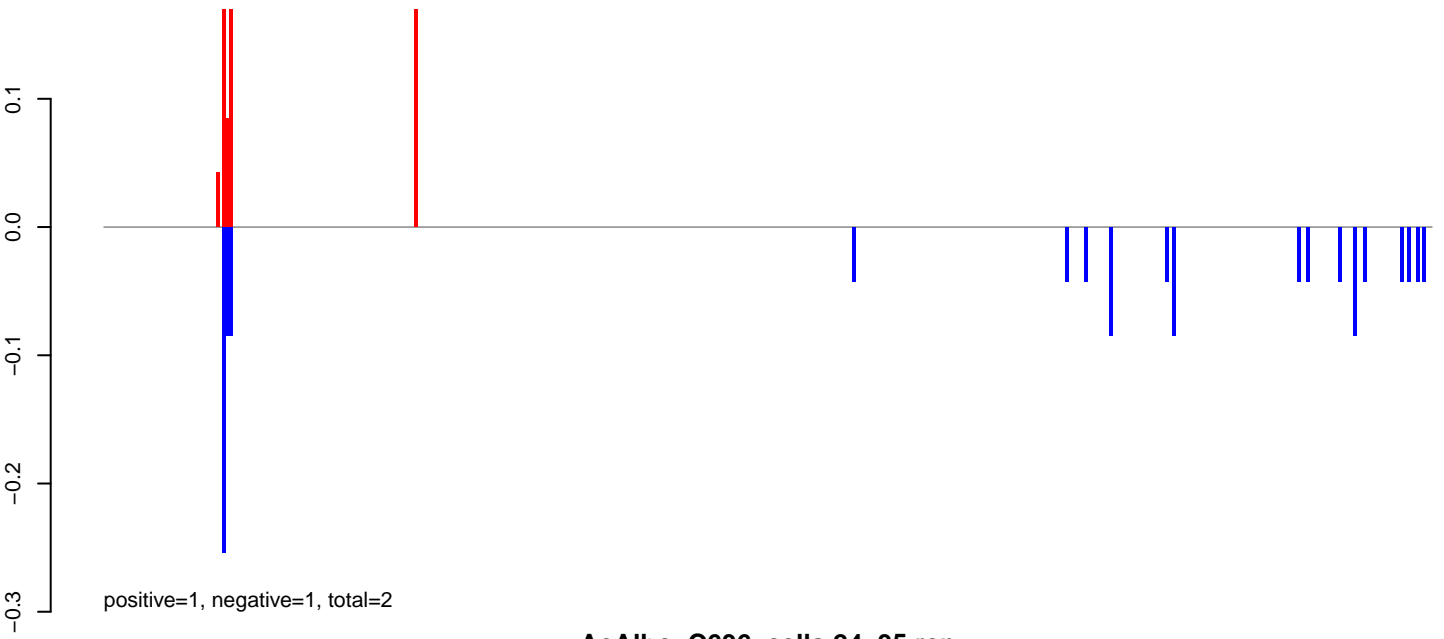


AeAlbo_C636_cells.rep

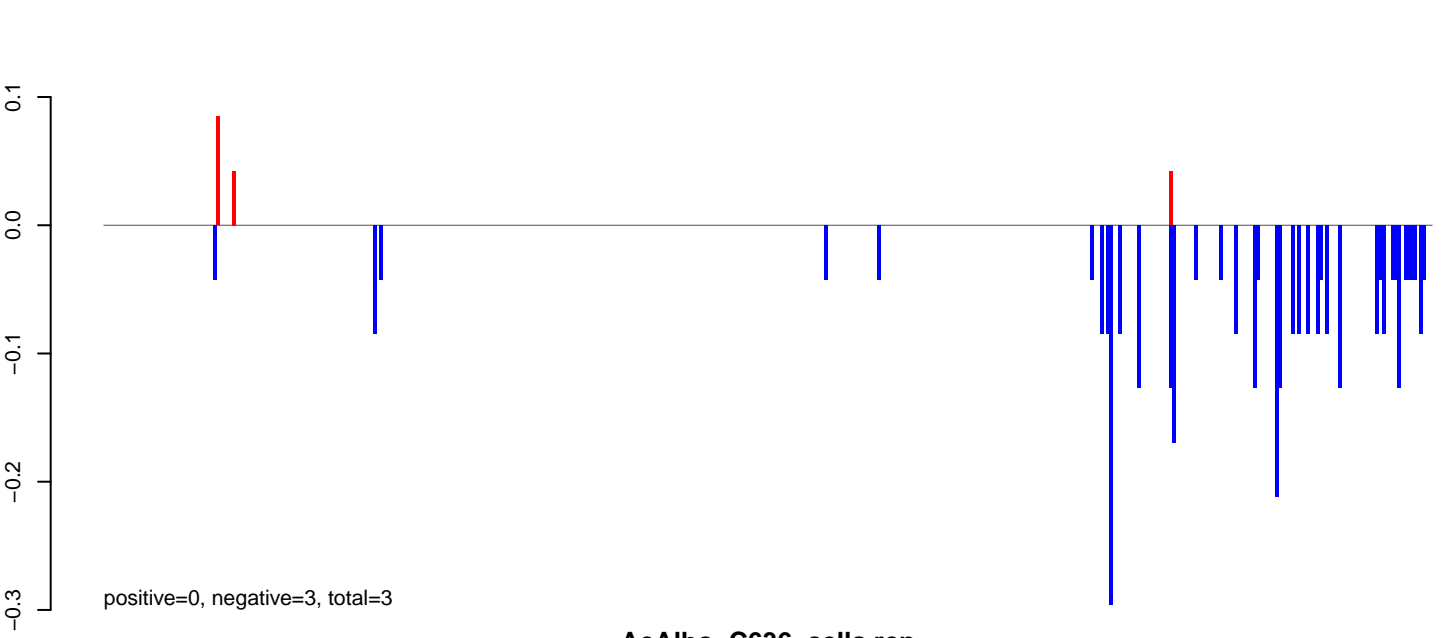


0 1000 2000 3000 4000

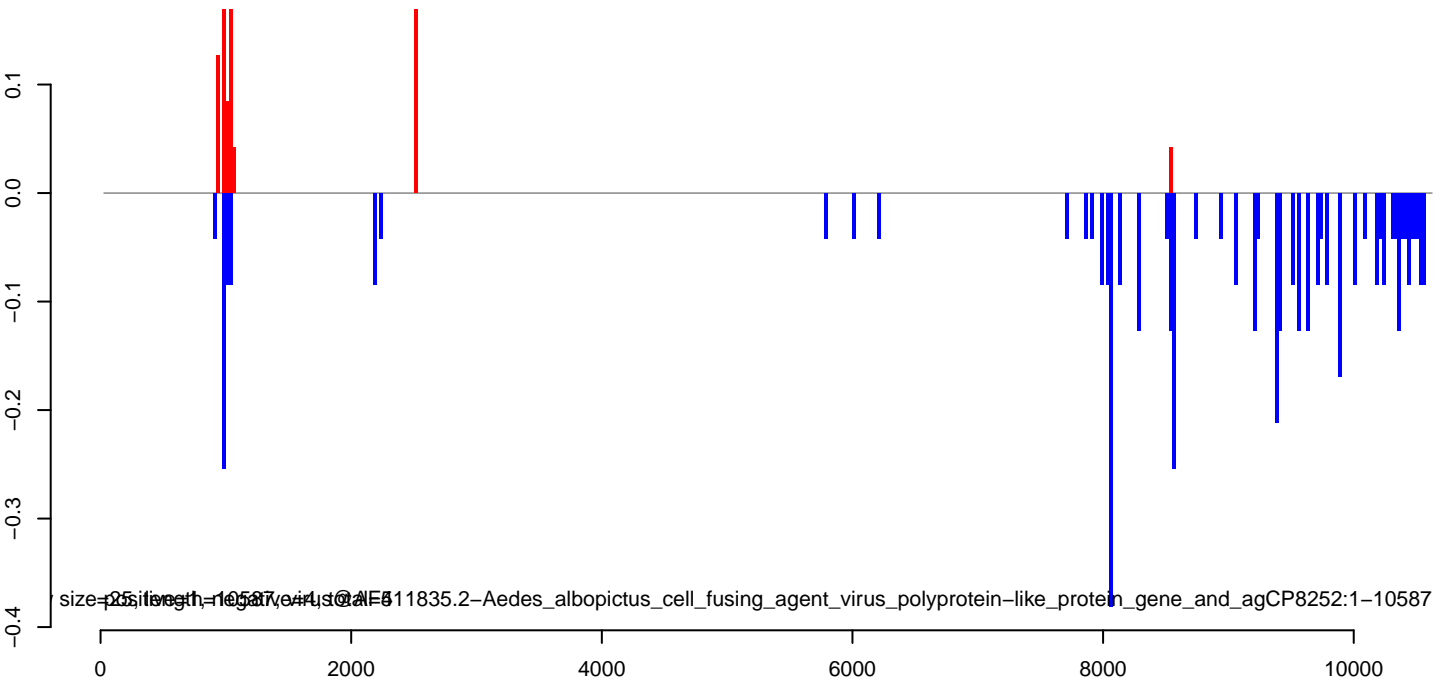
AeAlbo_C636_cells.18_23.rep



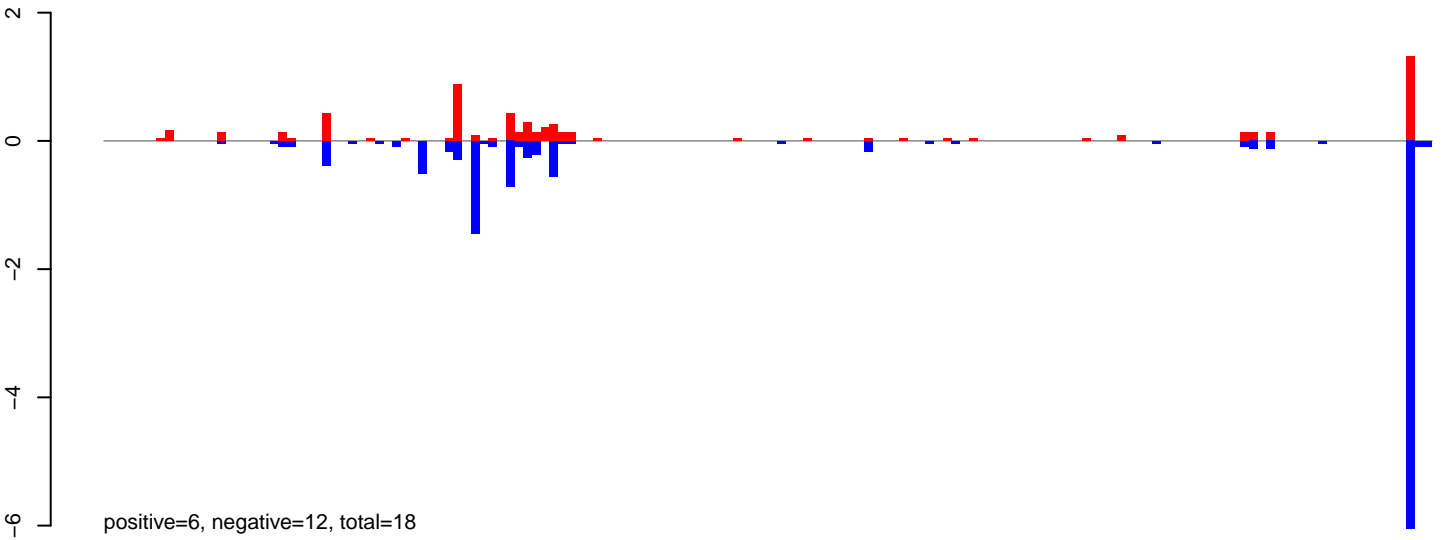
AeAlbo_C636_cells.24_35.rep



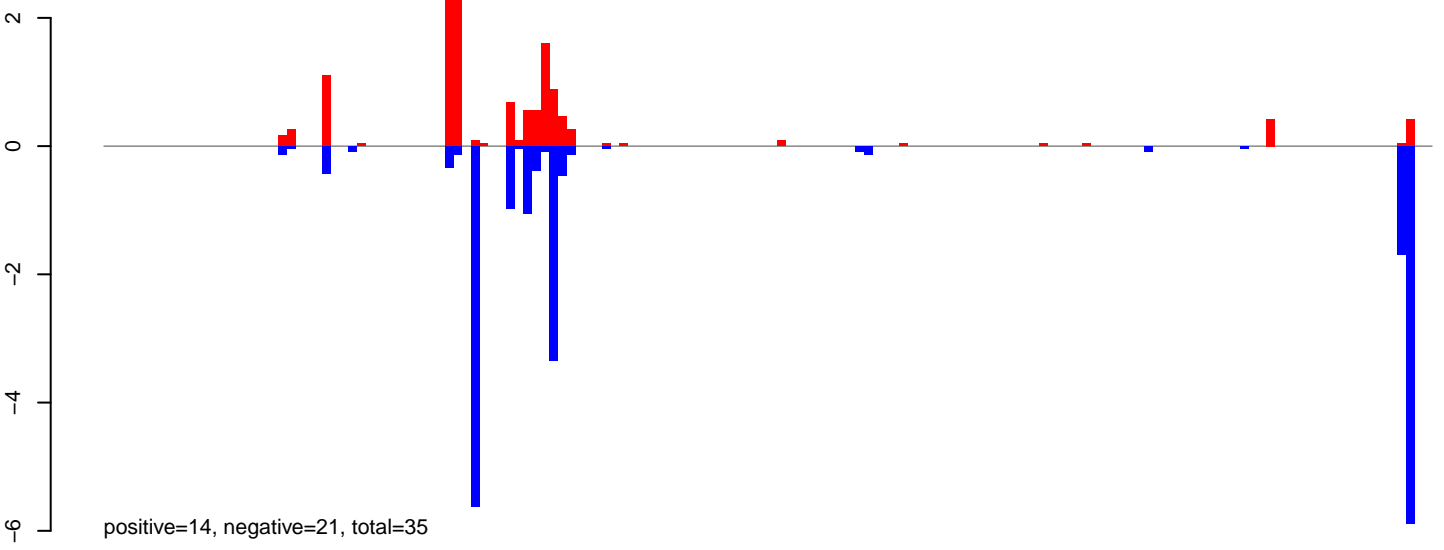
AeAlbo_C636_cells.rep



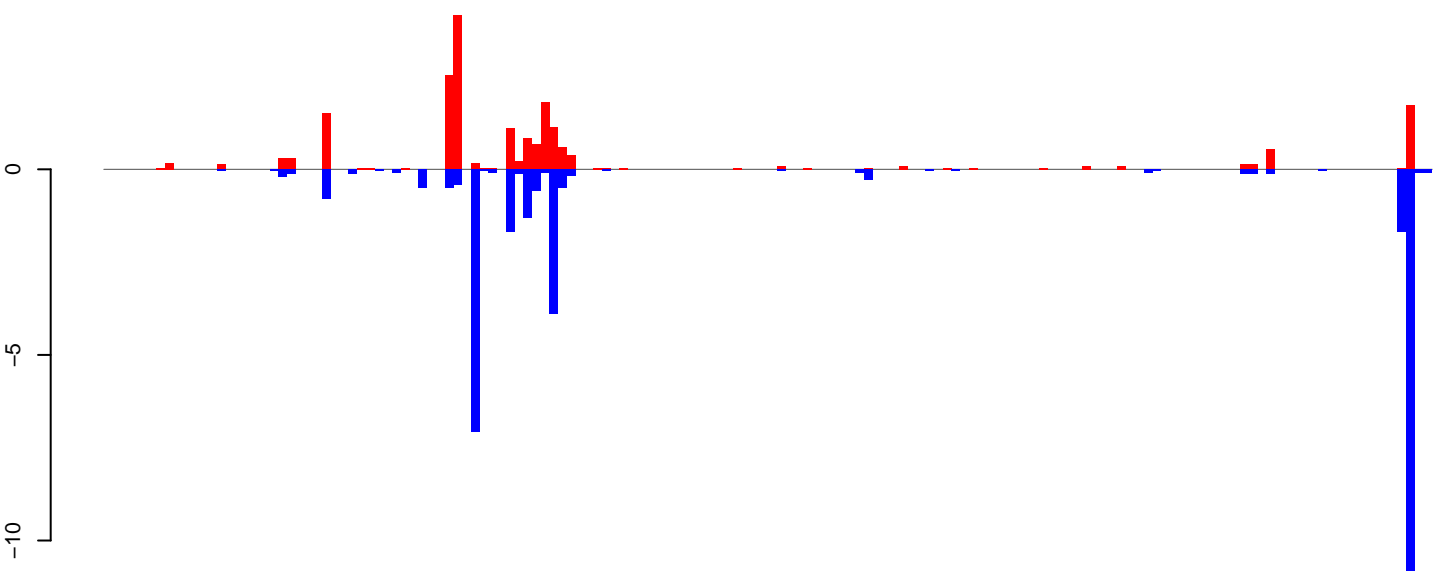
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



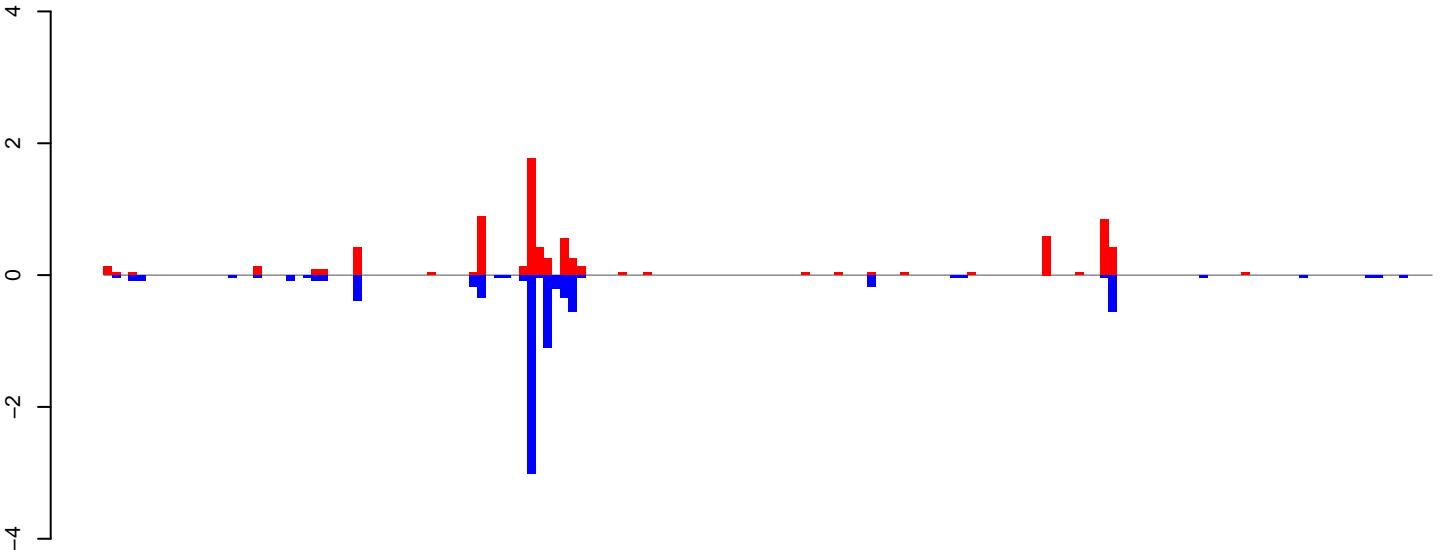
AeAlbo_C636_cells.rep



Window size=25, length=3790, virus@AeAeg_Thai_Densovirus_c30446_g1_i1_len=3790:1-3790

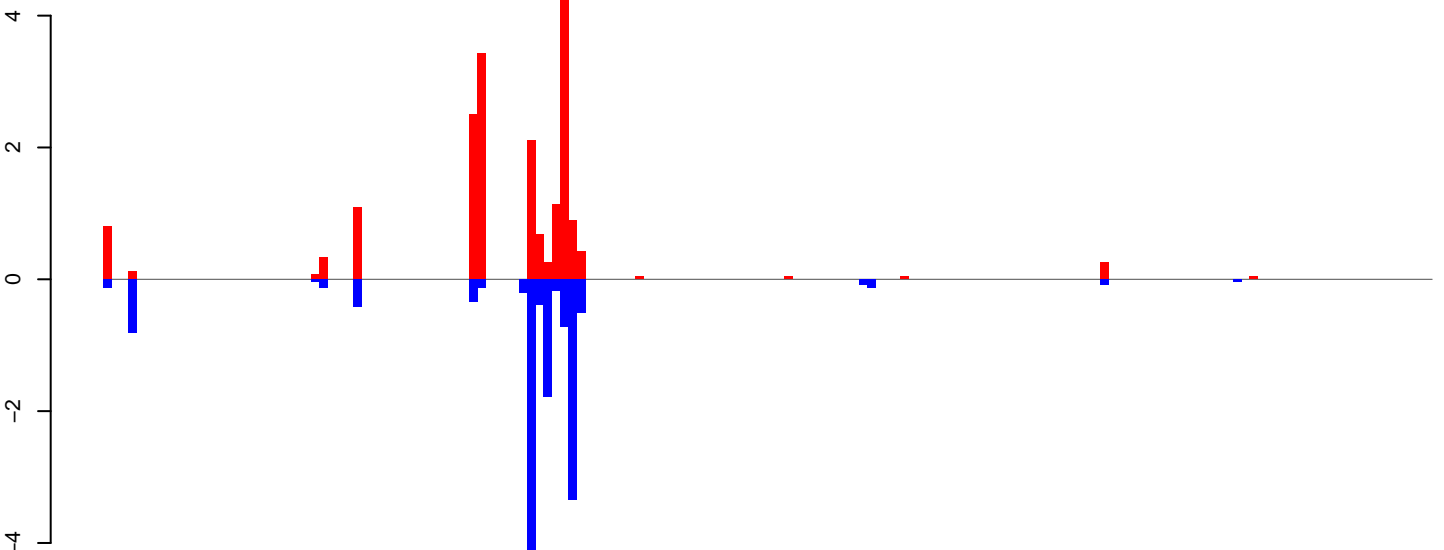
0 1000 2000 3000

AeAlbo_C636_cells.18_23.rep



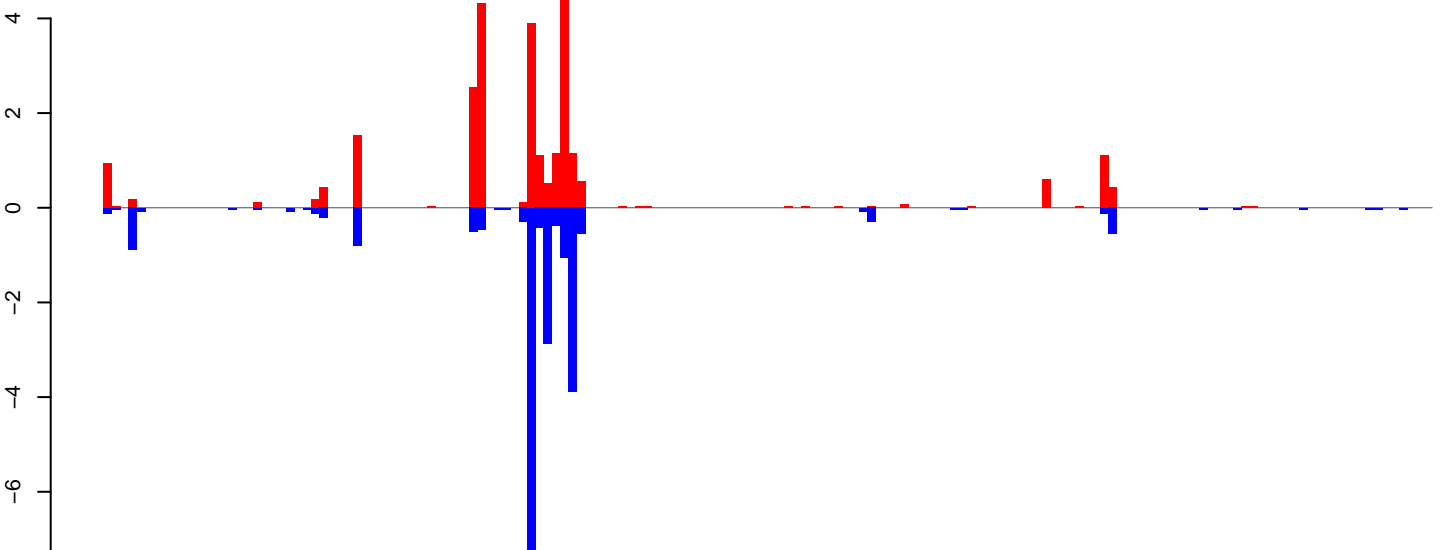
positive=8, negative=8, total=16

AeAlbo_C636_cells.24_35.rep



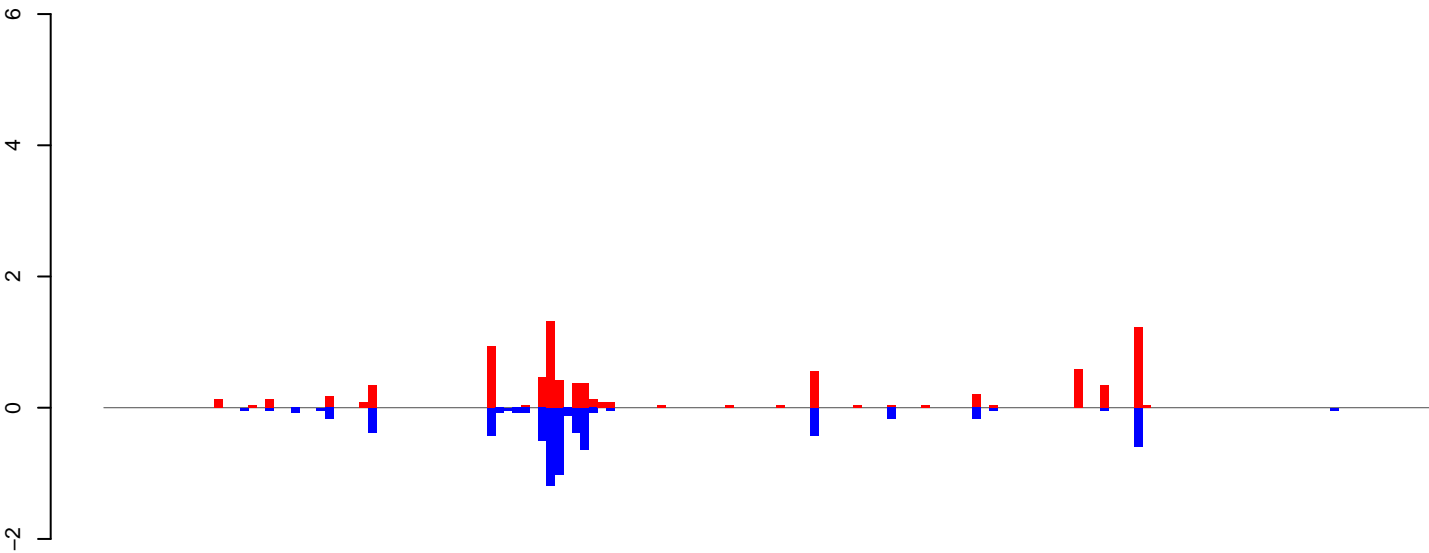
positive=19, negative=14, total=33

AeAlbo_C636_cells.rep



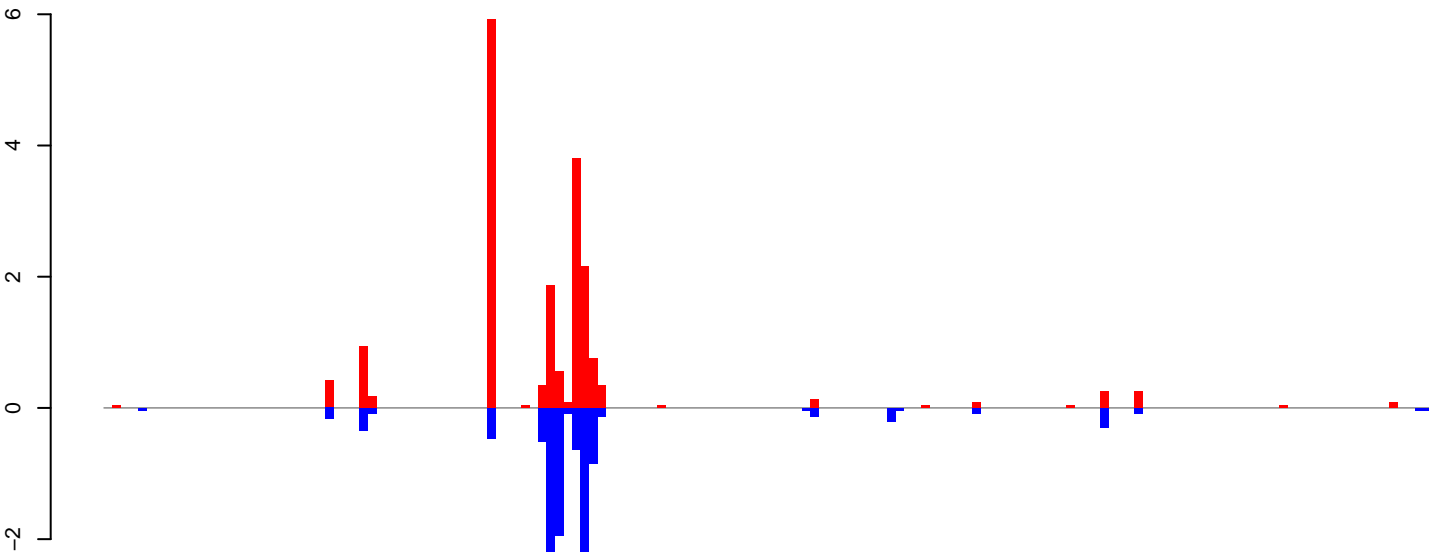
positive=26, negative=22, total=48 window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

AeAlbo_C636_cells.18_23.rep



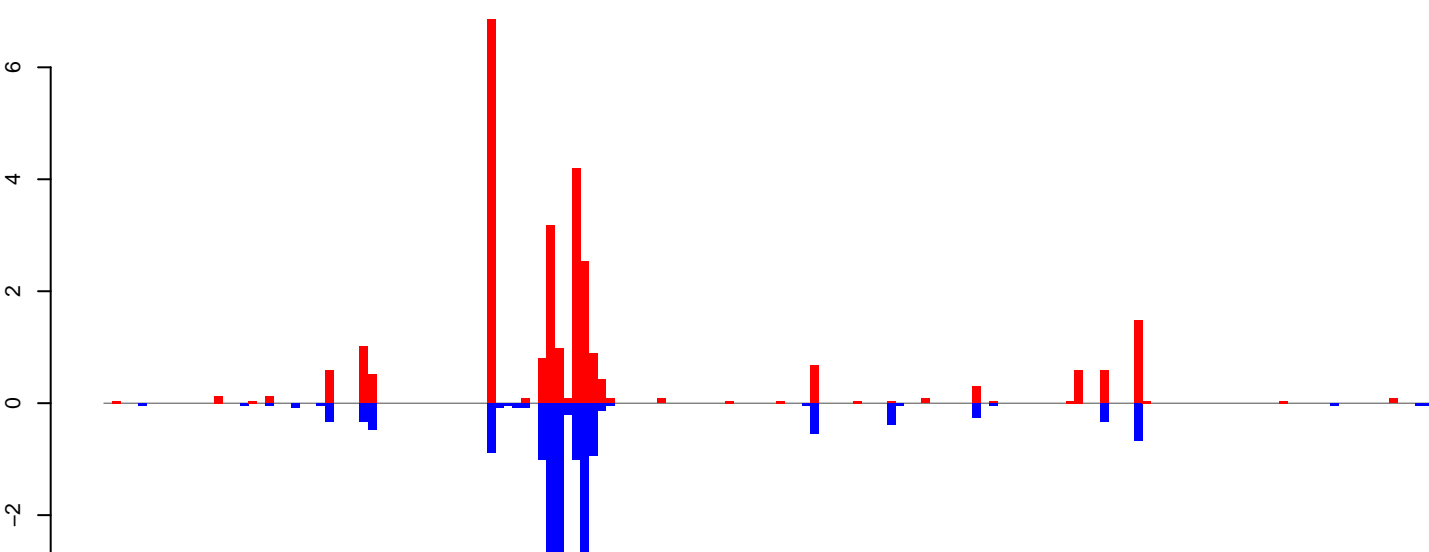
positive=8, negative=7, total=15

AeAlbo_C636_cells.24_35.rep



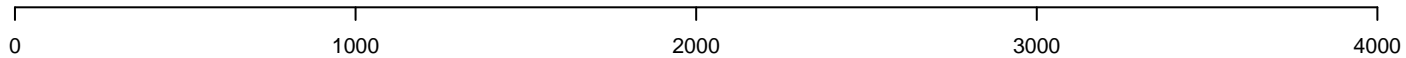
positive=18, negative=12, total=30

AeAlbo_C636_cells.rep

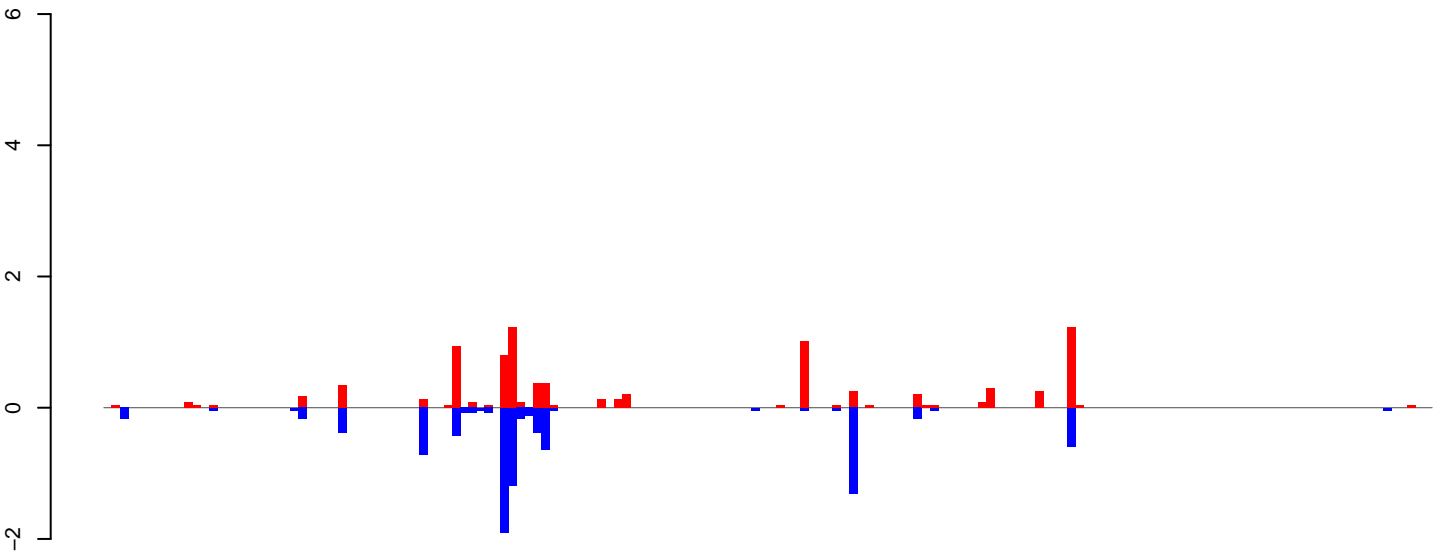


positive=27, negative=19, total=46

Window size=25, length=3893, virus@GU452720.1-Mosquito_densovirus_BR_07:1-3893

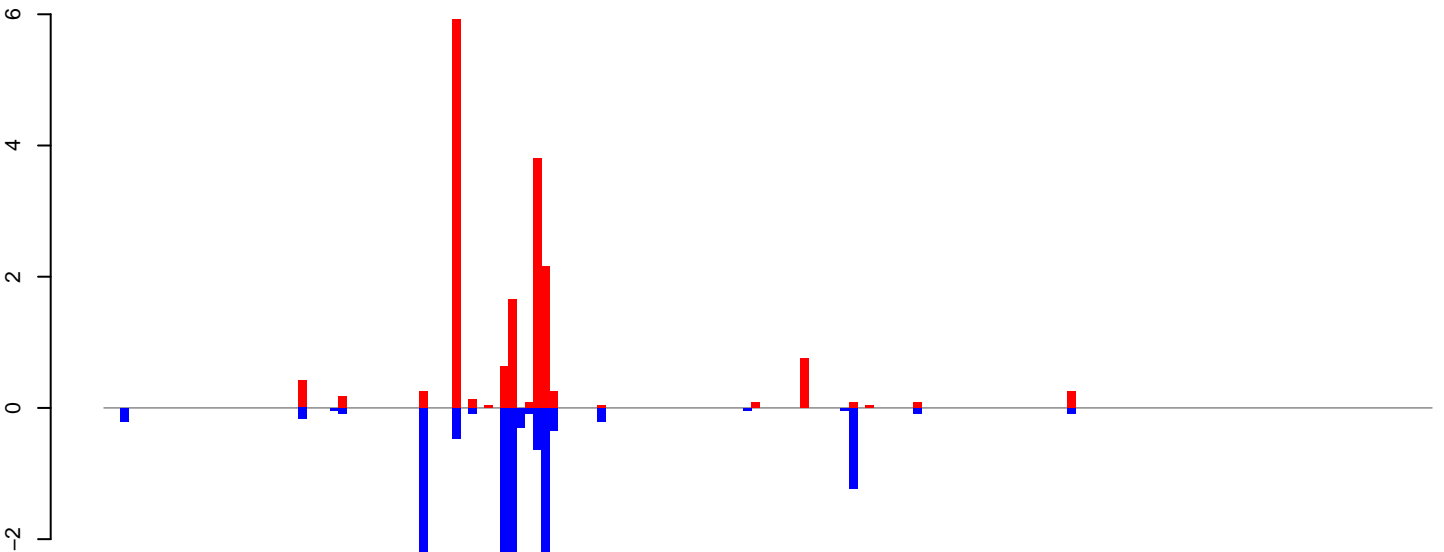


AeAlbo_C636_cells.18_23.rep



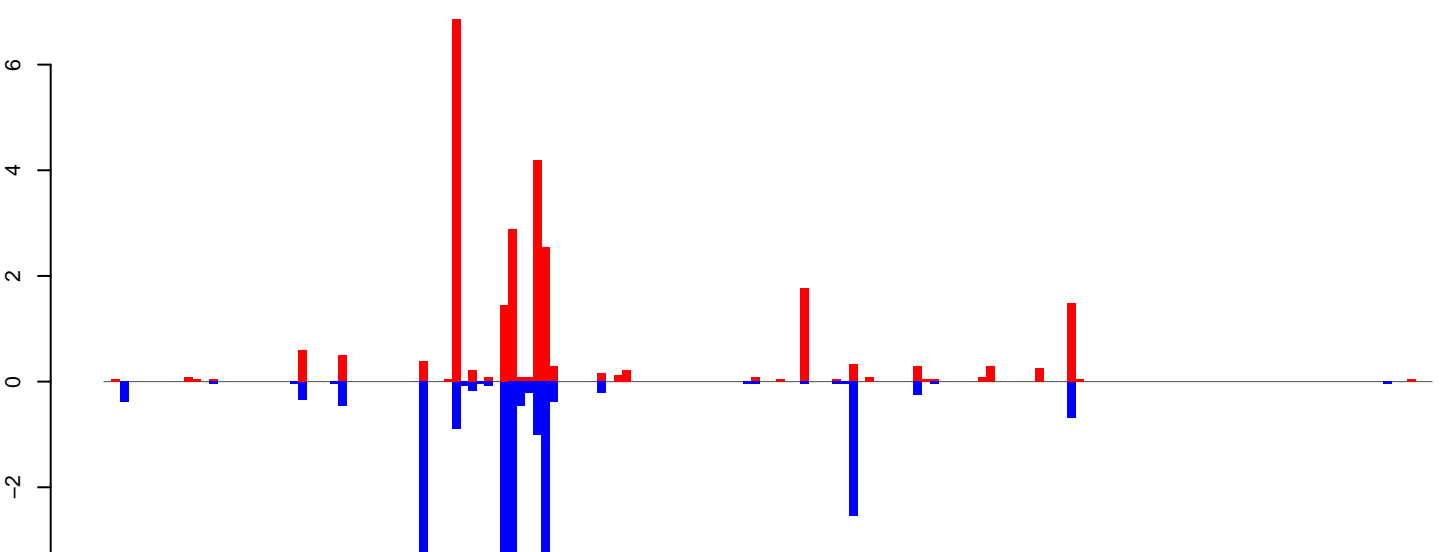
positive=9, negative=9, total=18

AeAlbo_C636_cells.24_35.rep



positive=17, negative=15, total=32

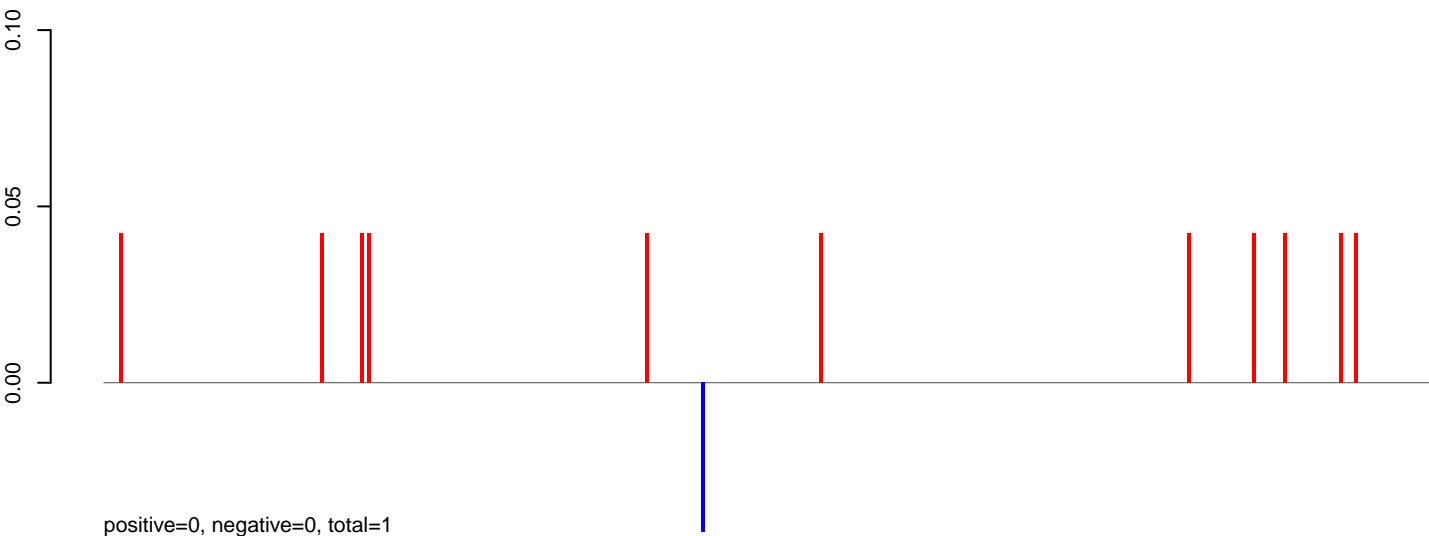
AeAlbo_C636_cells.rep



bow size=25, length=4094, negative=24, total=55
Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural:1-4094

0 1000 2000 3000 4000

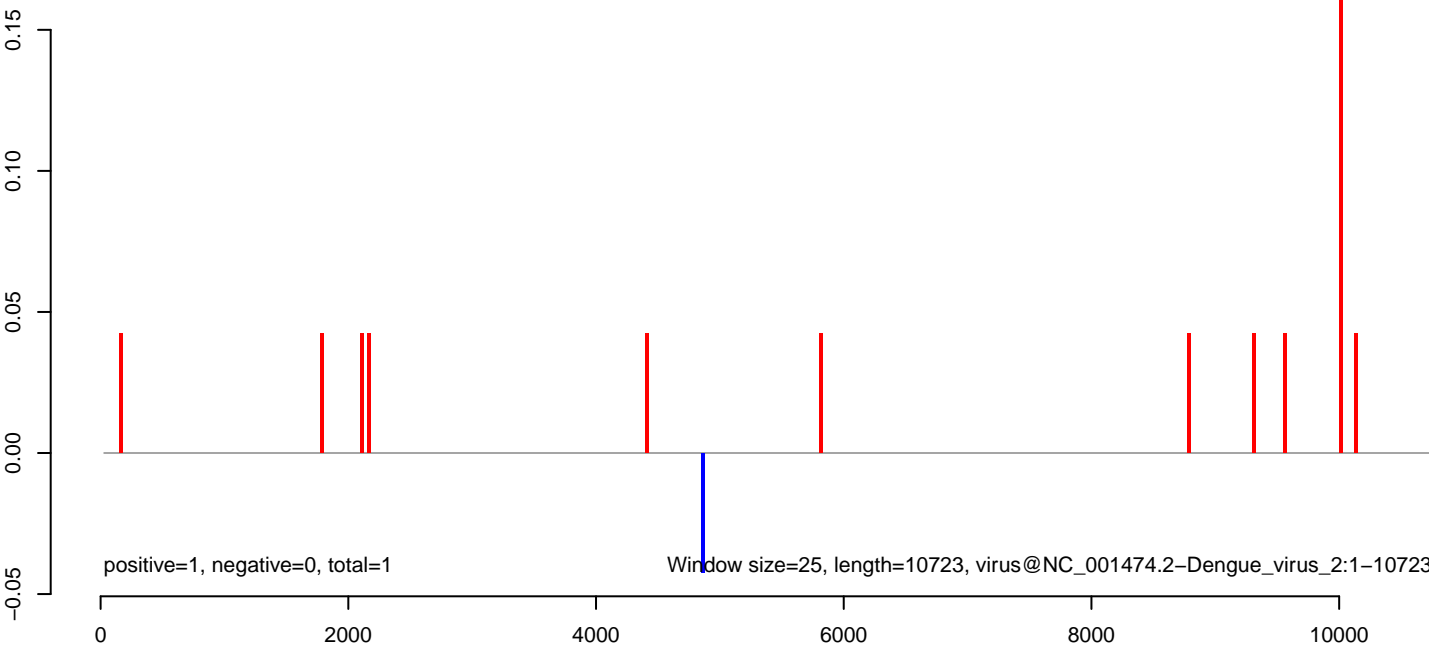
AeAlbo_C636_cells.18_23.rep



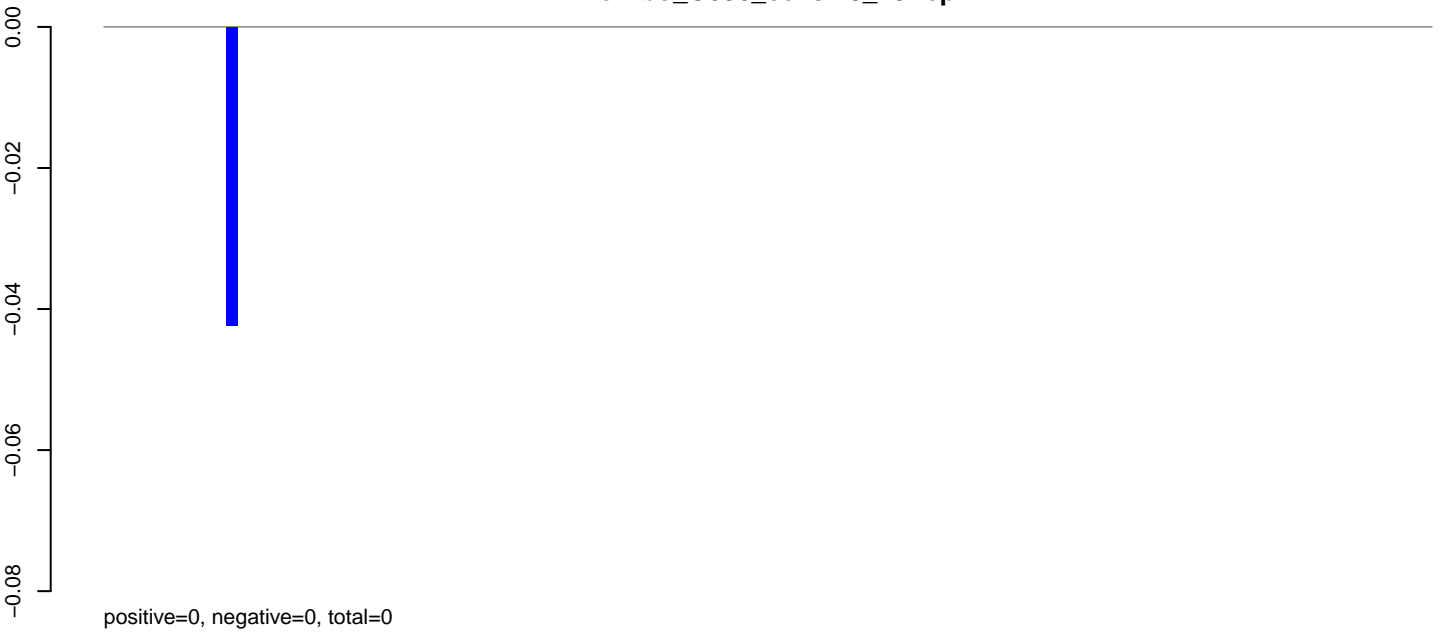
AeAlbo_C636_cells.24_35.rep



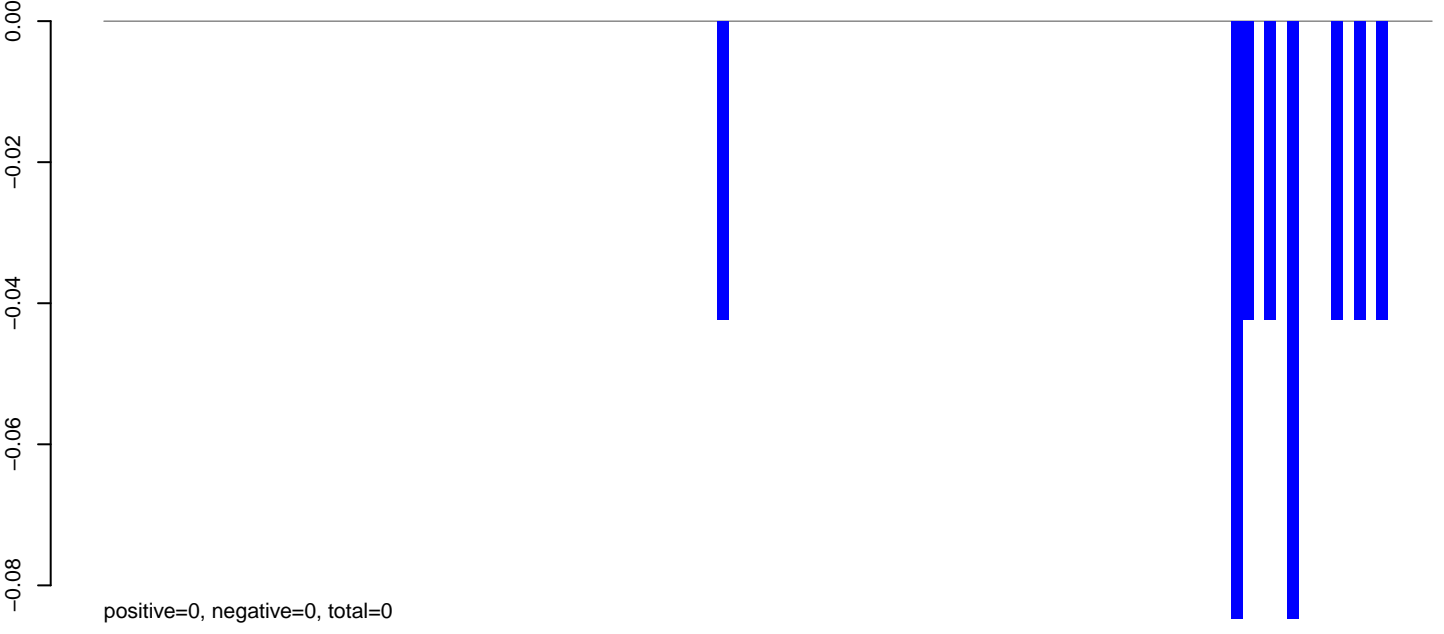
AeAlbo_C636_cells.rep



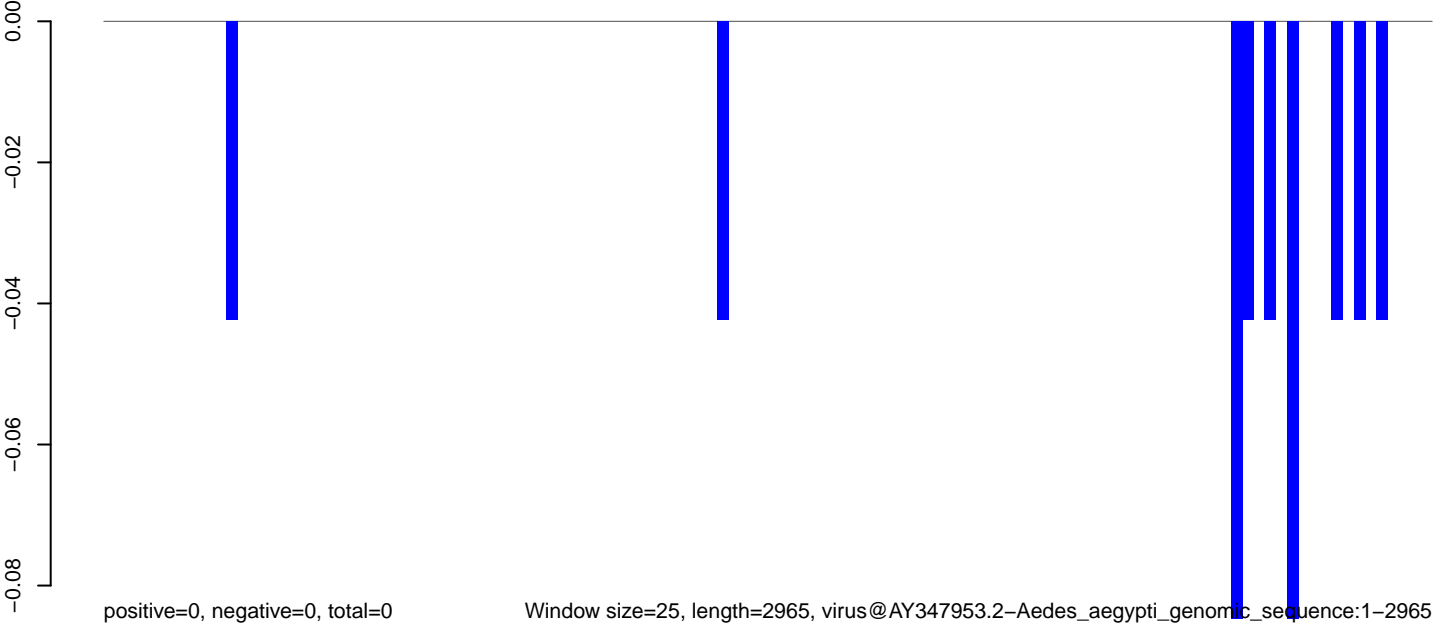
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



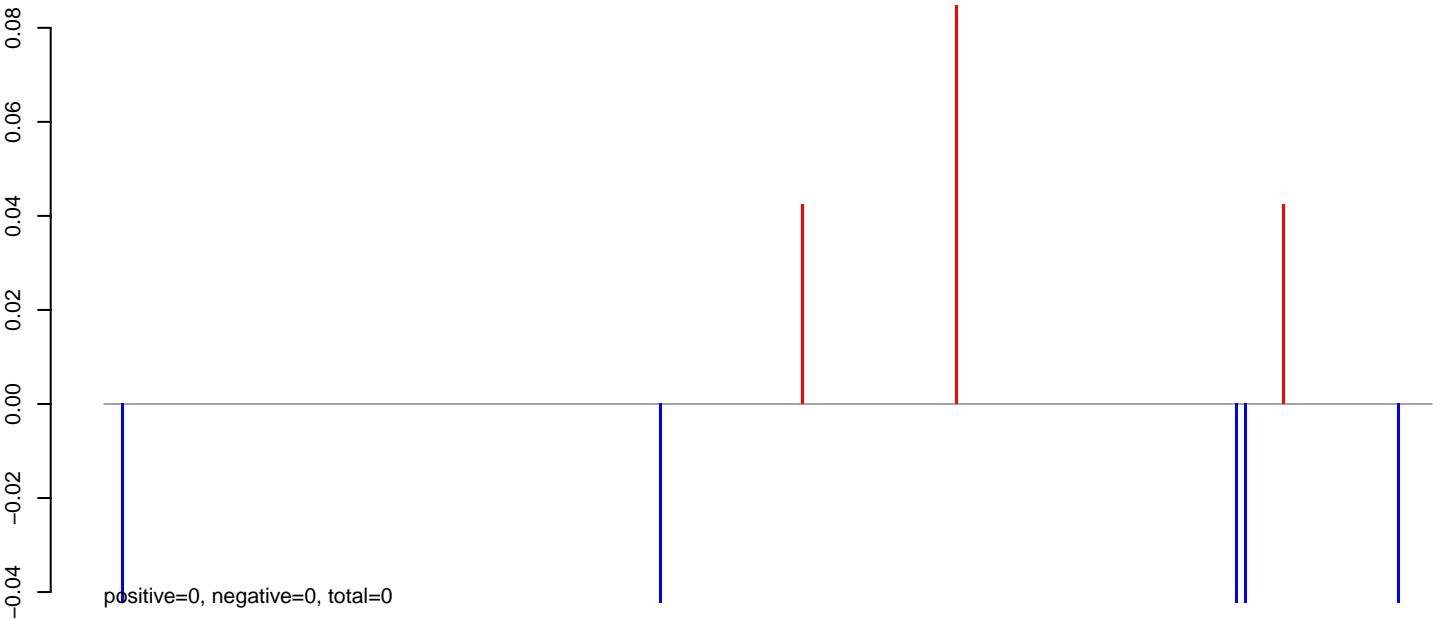
AeAlbo_C636_cells.rep



Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

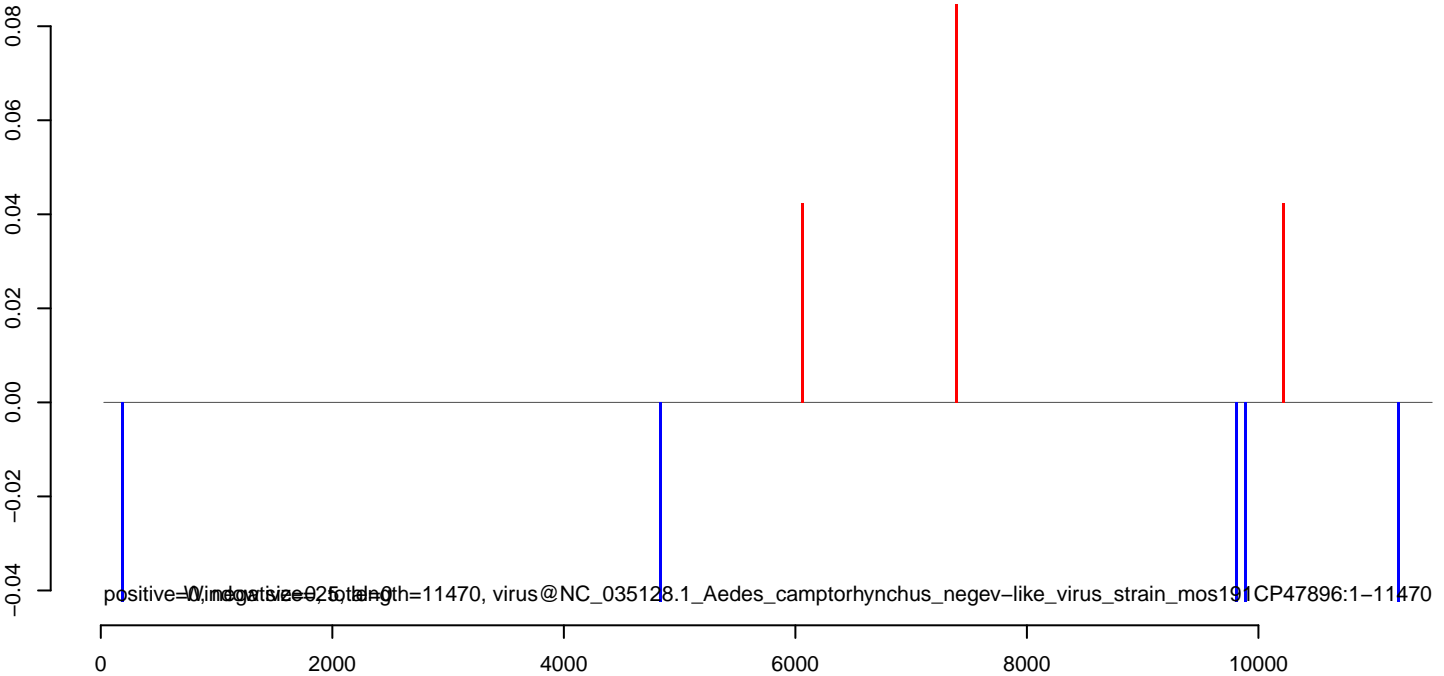
AeAlbo_C636_cells.18_23.rep



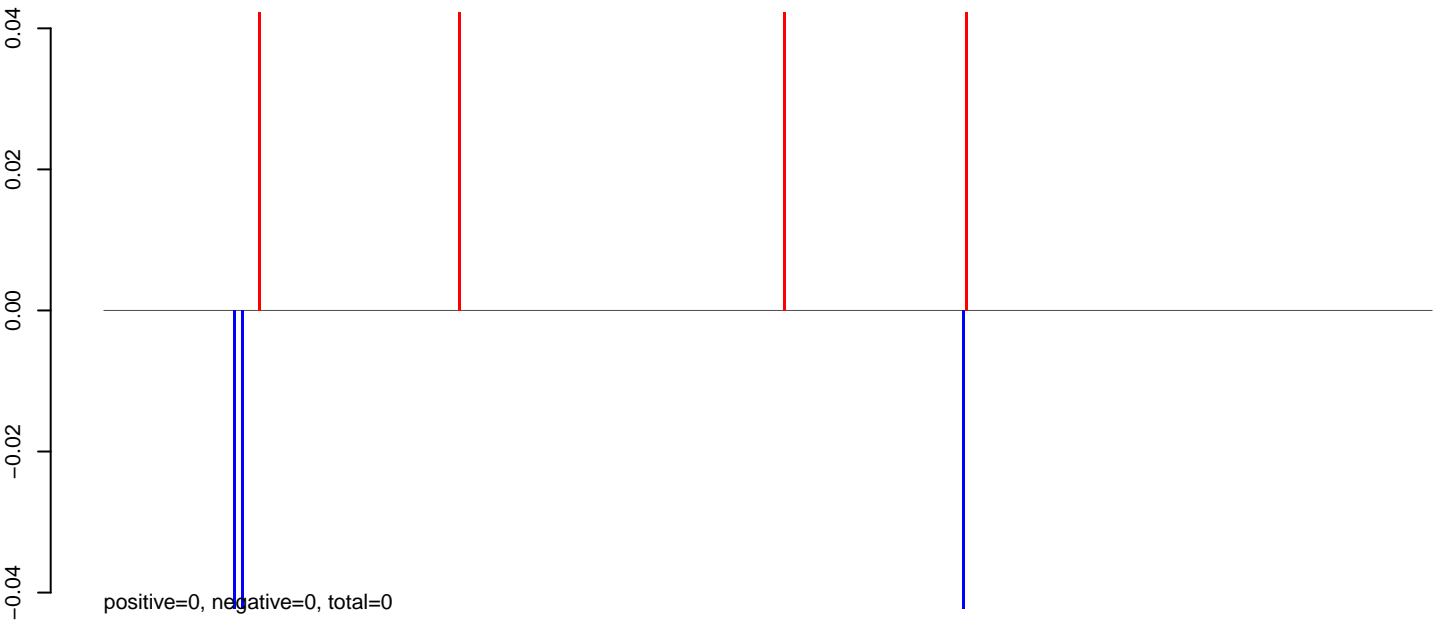
AeAlbo_C636_cells.24_35.rep



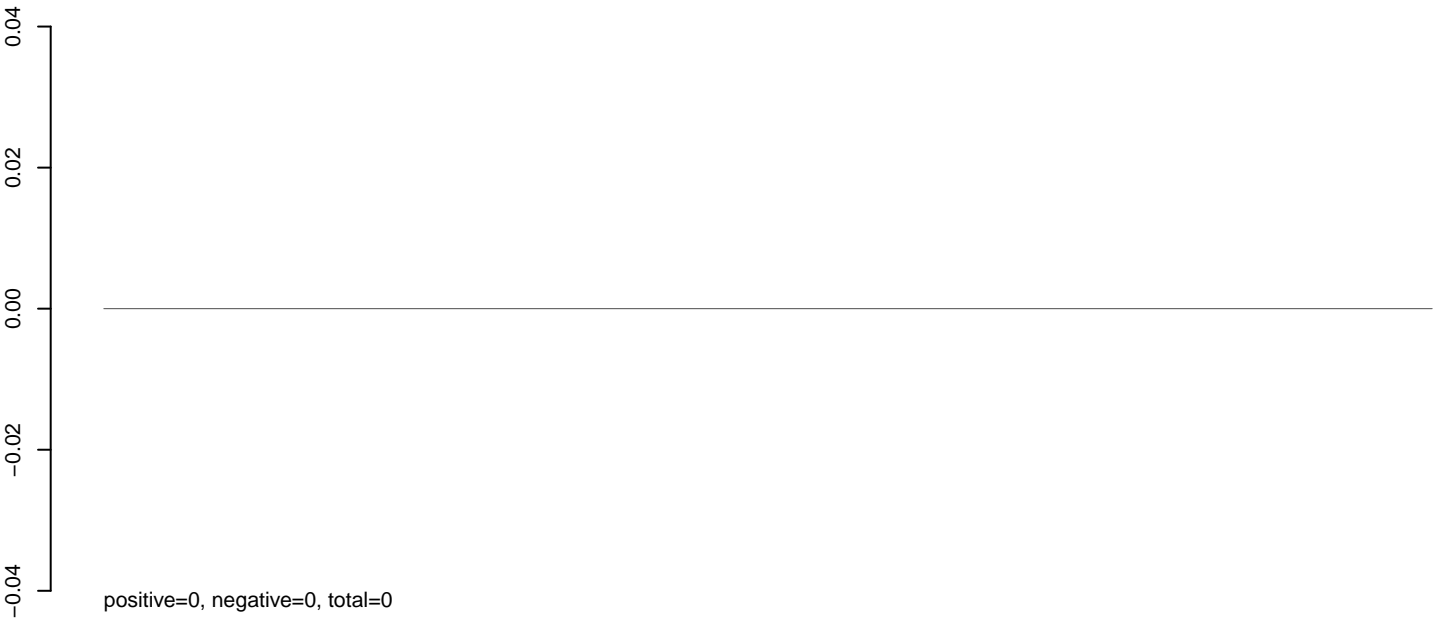
AeAlbo_C636_cells.rep



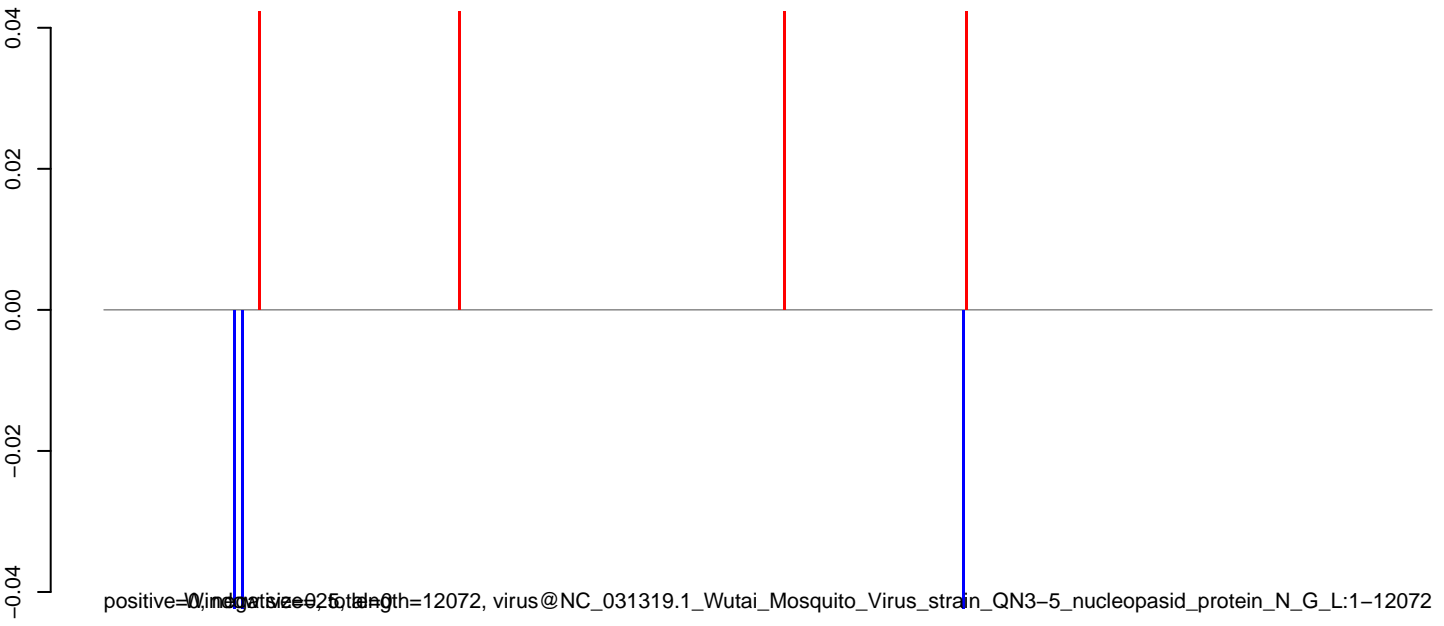
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

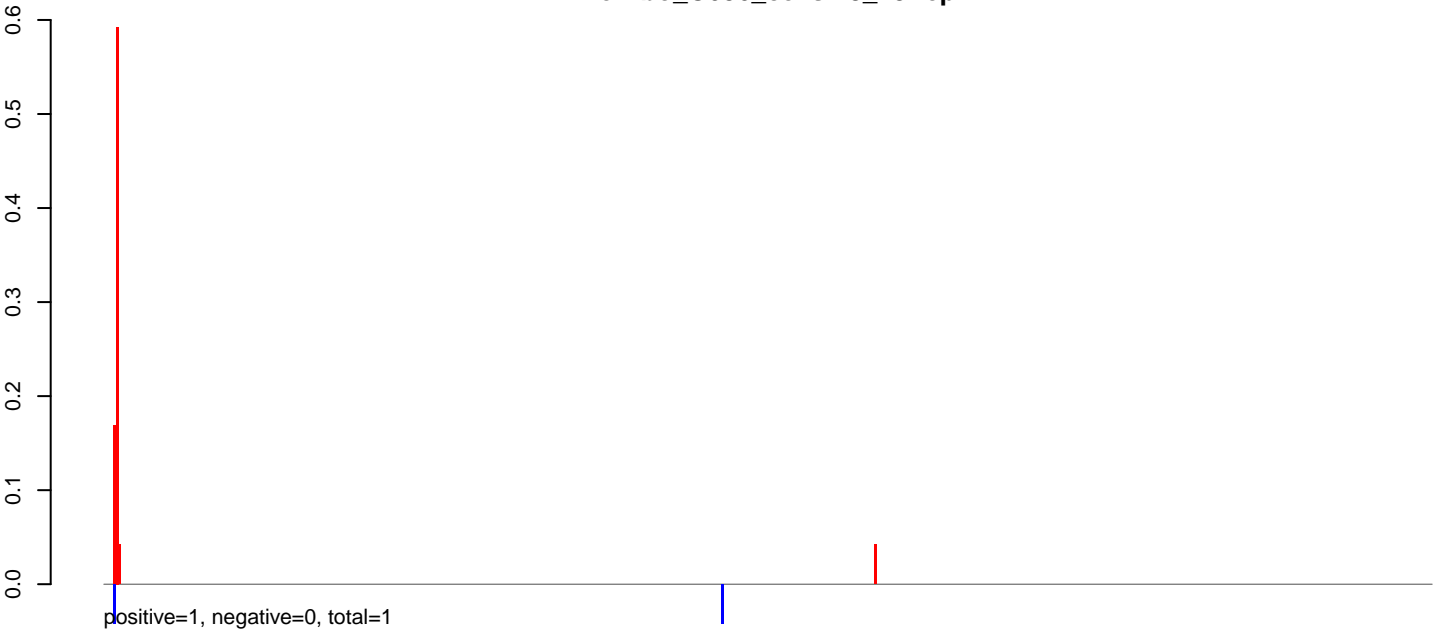


AeAlbo_C636_cells.rep

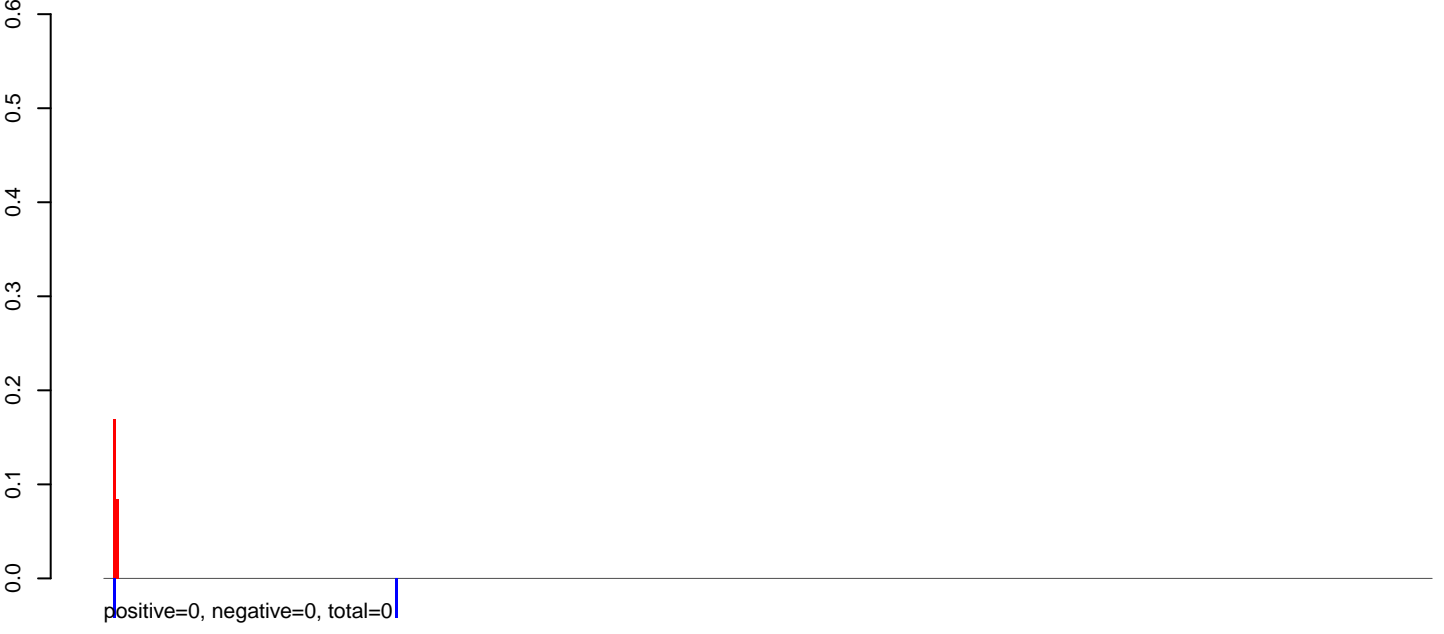


0 2000 4000 6000 8000 10000 12000
positive=0, negative=0, total=0, virus@NC_031319.1_Wutai_Mosquito_Virus_strain_QN3-5_nucleopasid_protein_N_G_L:1-12072

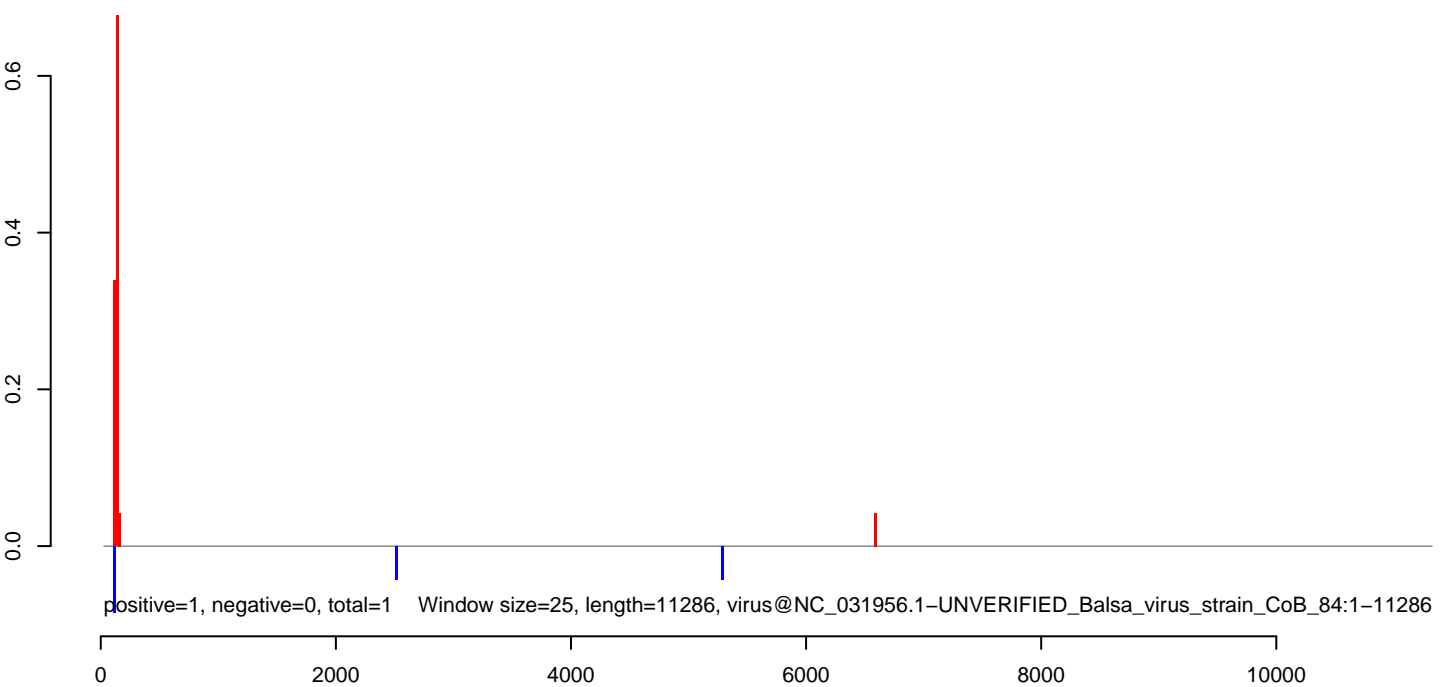
AeAlbo_C636_cells.18_23.rep



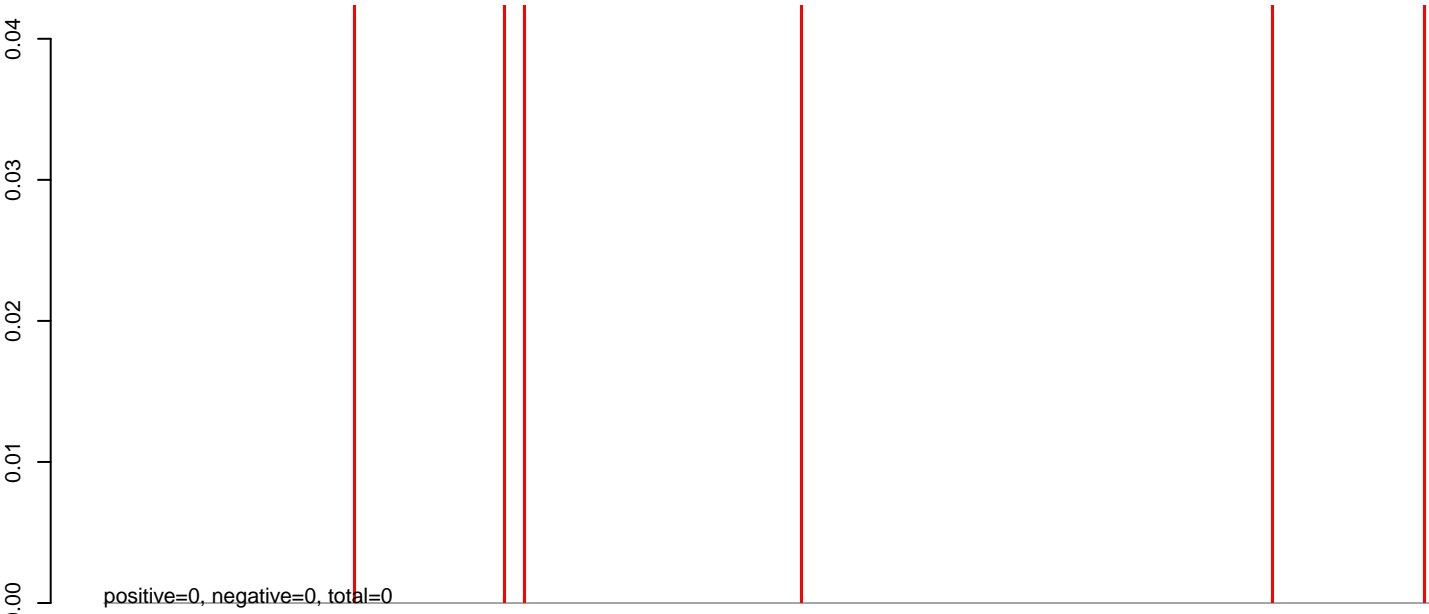
AeAlbo_C636_cells.24_35.rep



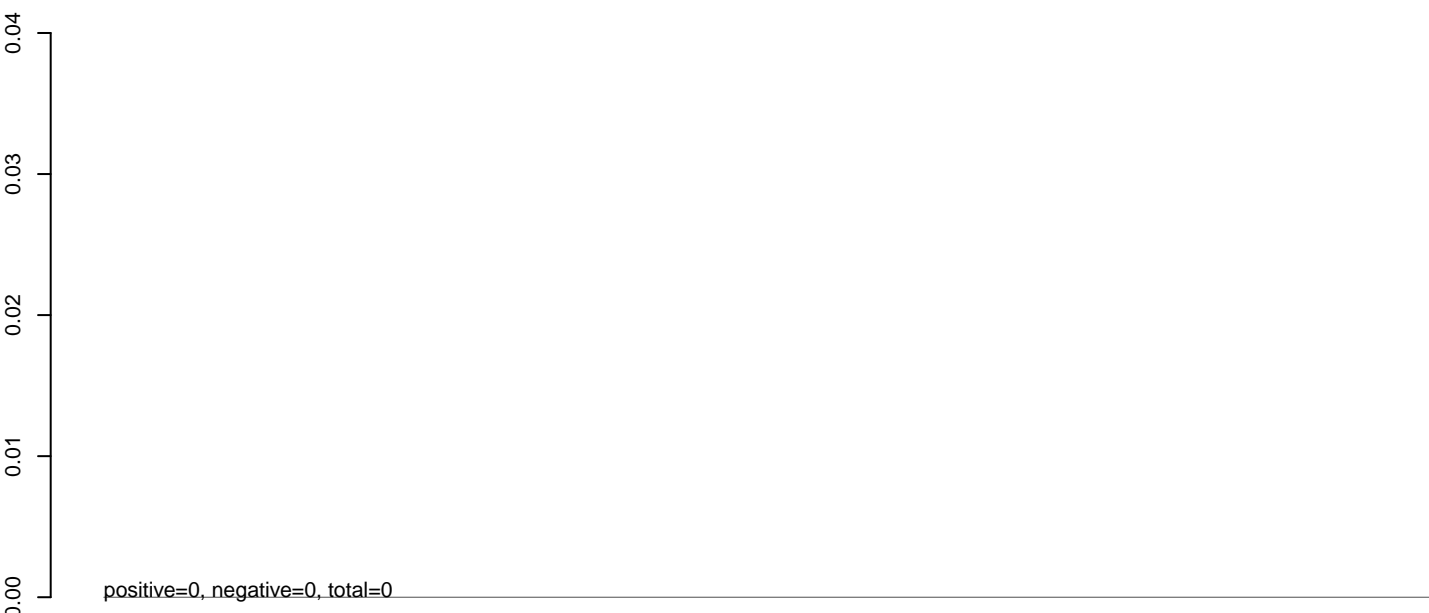
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



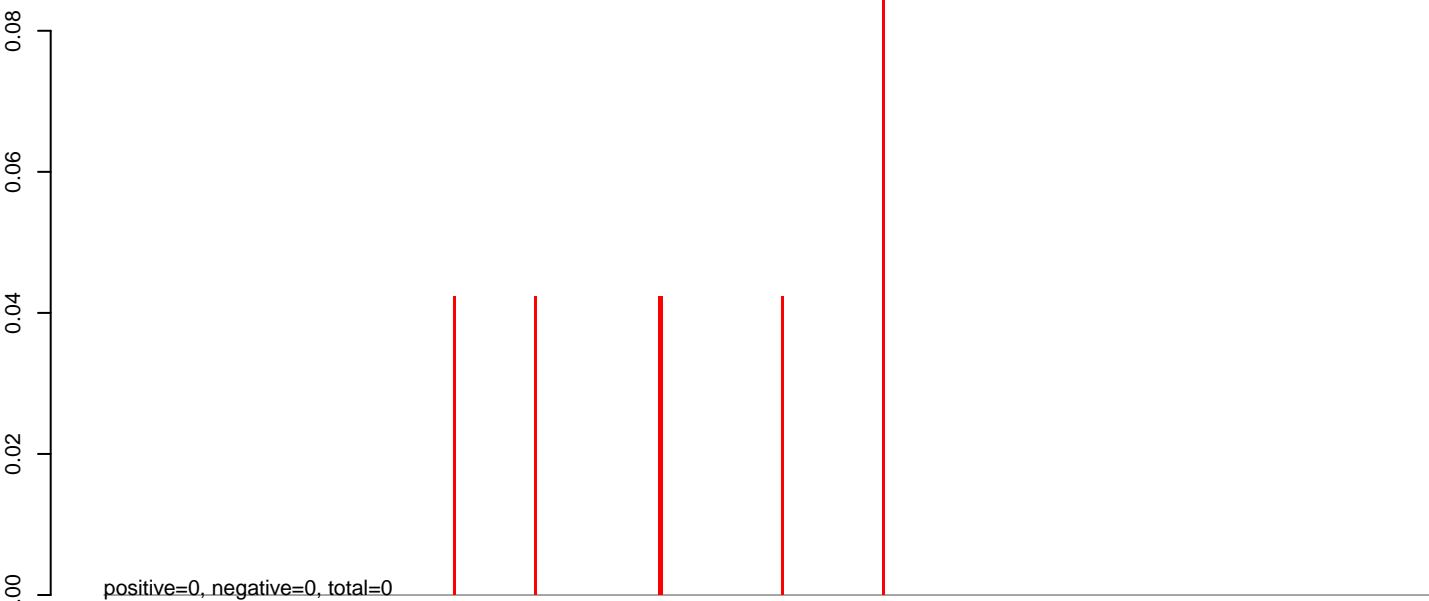
AeAlbo_C636_cells.24_35.rep



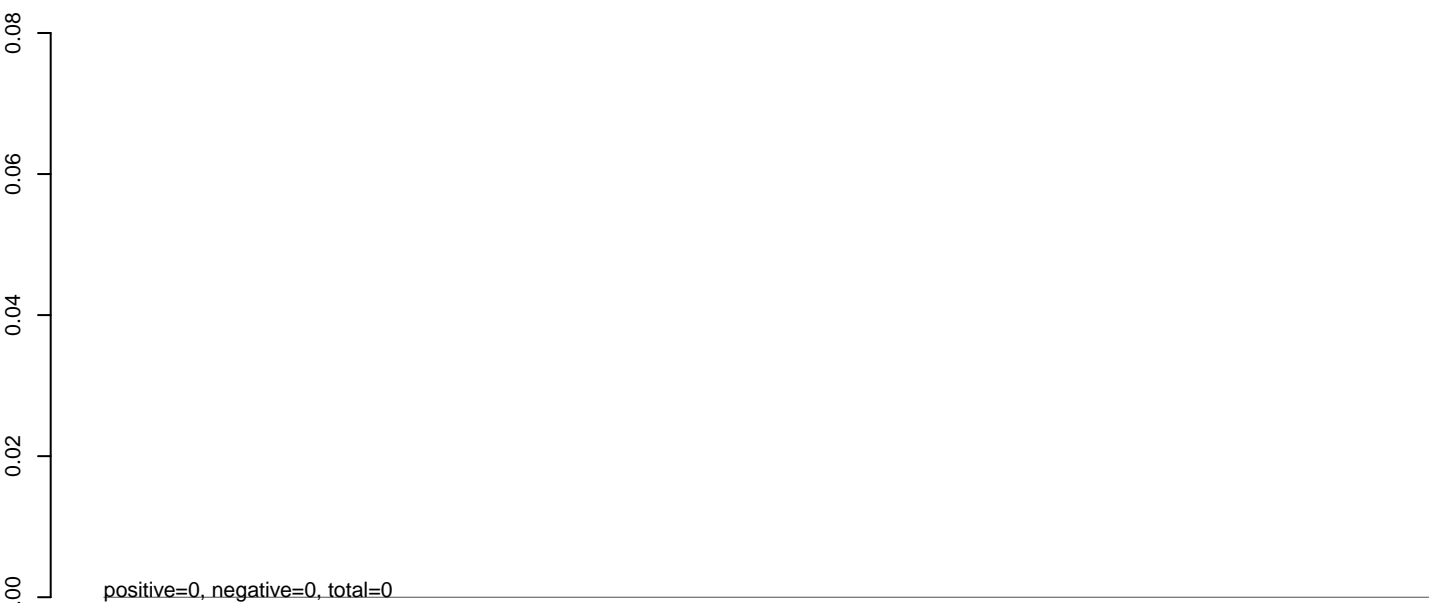
AeAlbo_C636_cells.rep



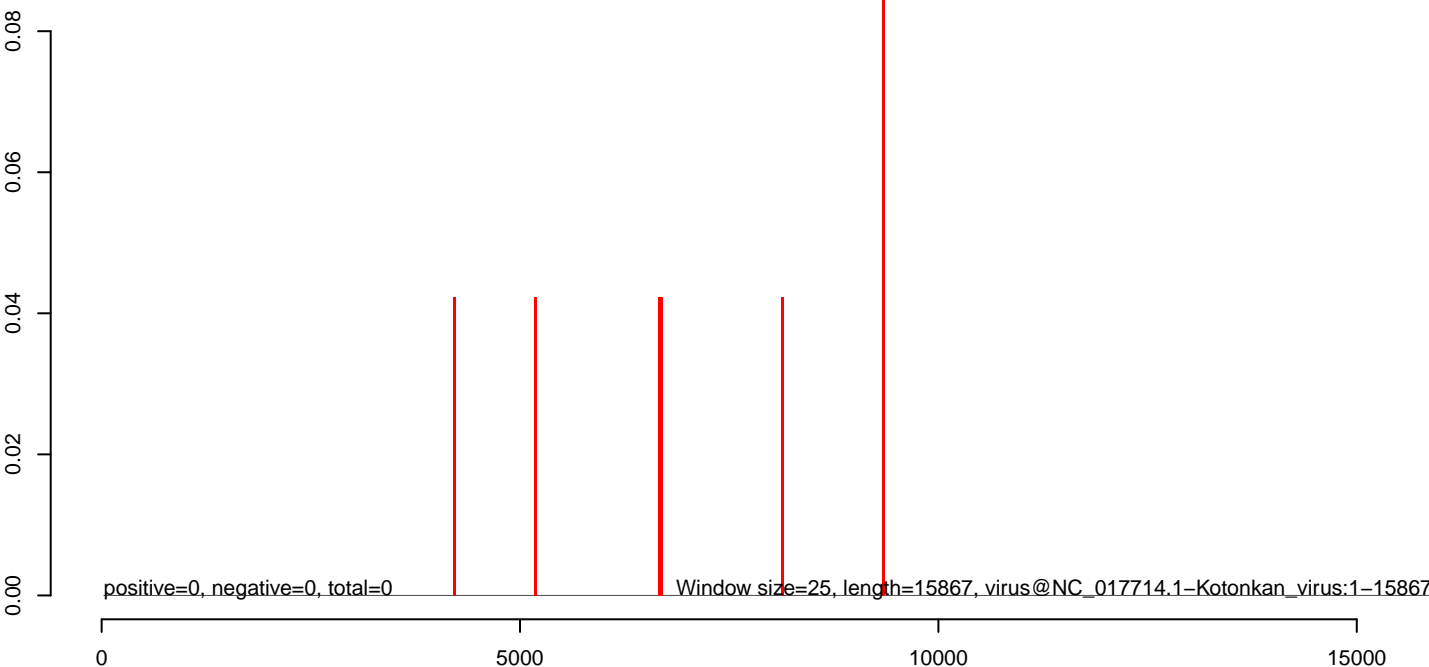
AeAlbo_C636_cells.18_23.rep



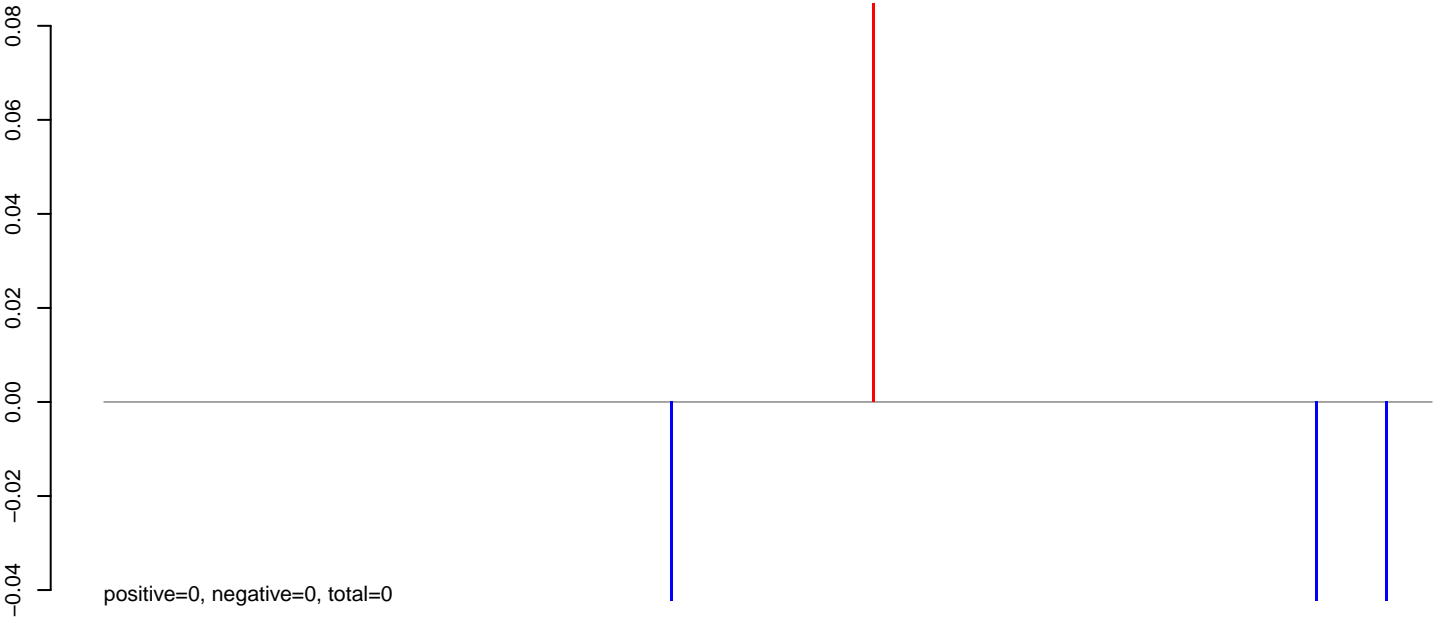
AeAlbo_C636_cells.24_35.rep



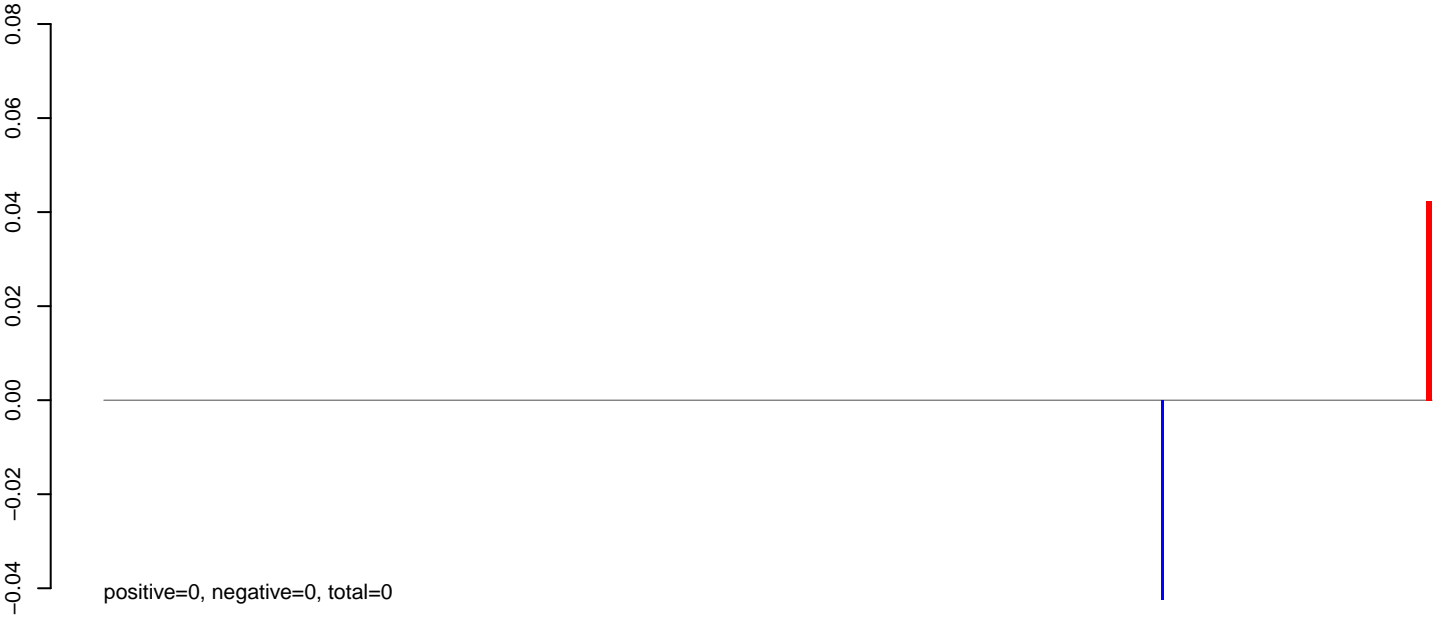
AeAlbo_C636_cells.rep



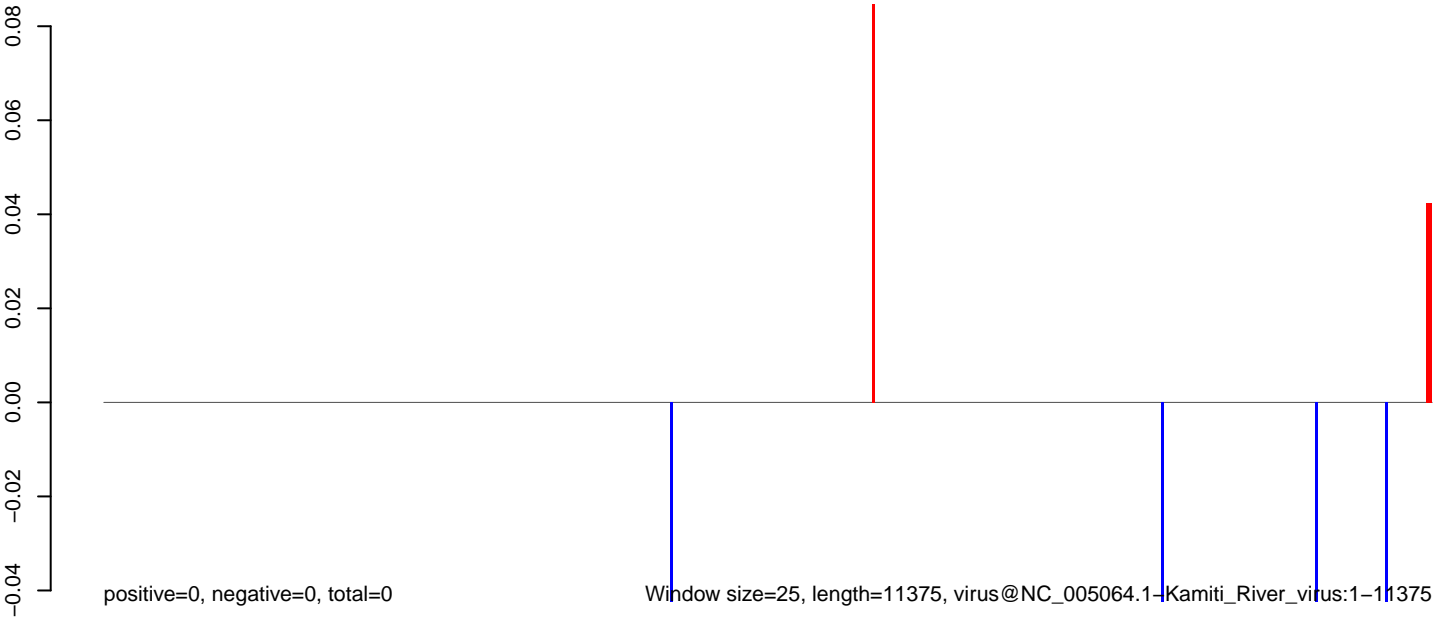
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

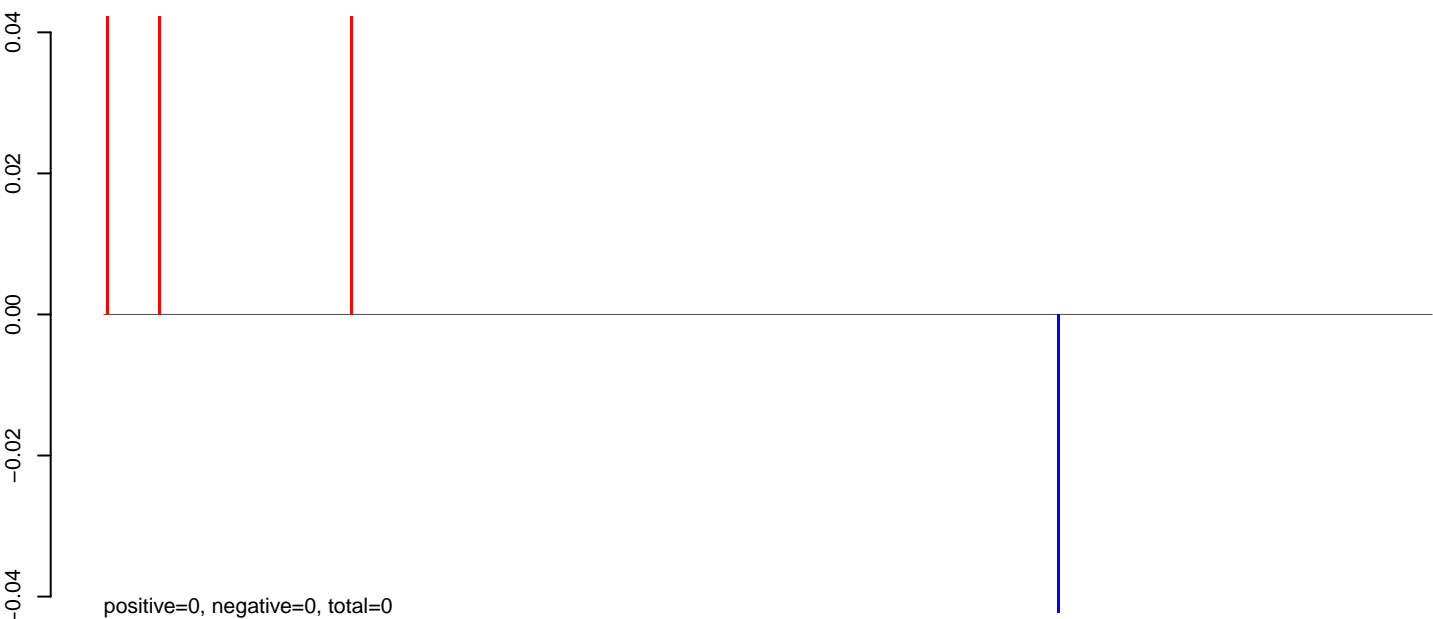


AeAlbo_C636_cells.rep

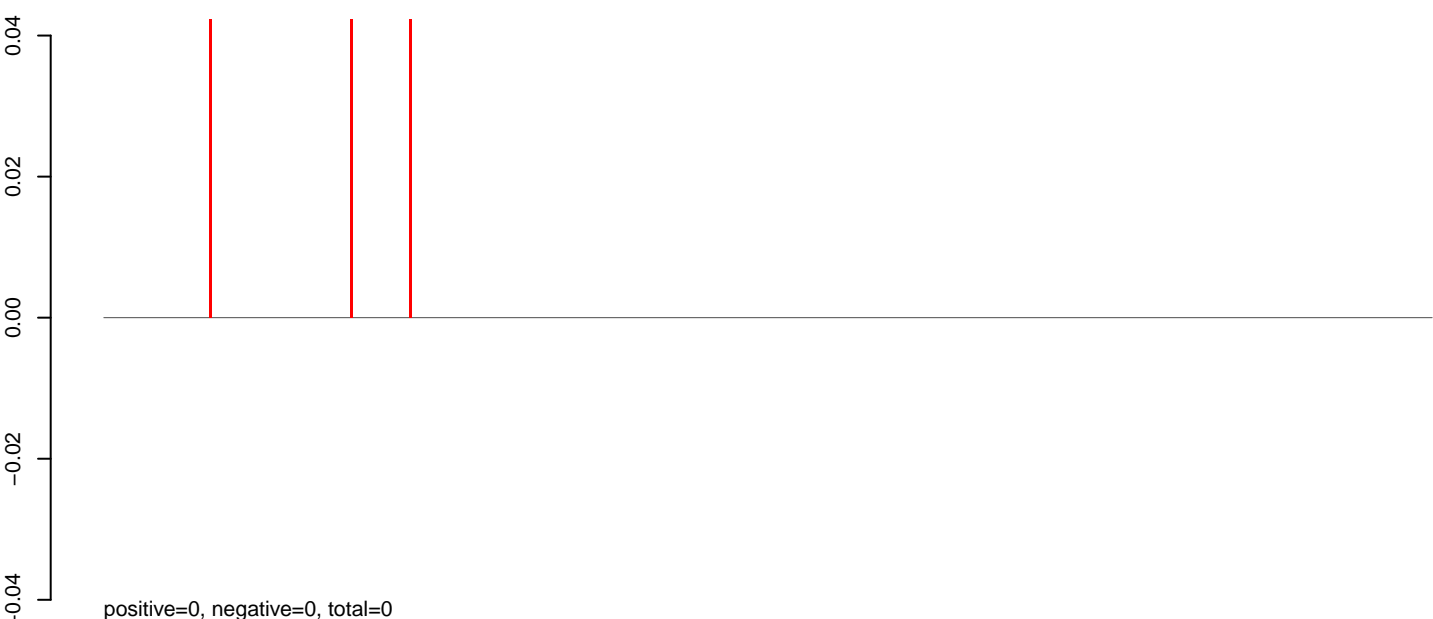


0 2000 4000 6000 8000 10000

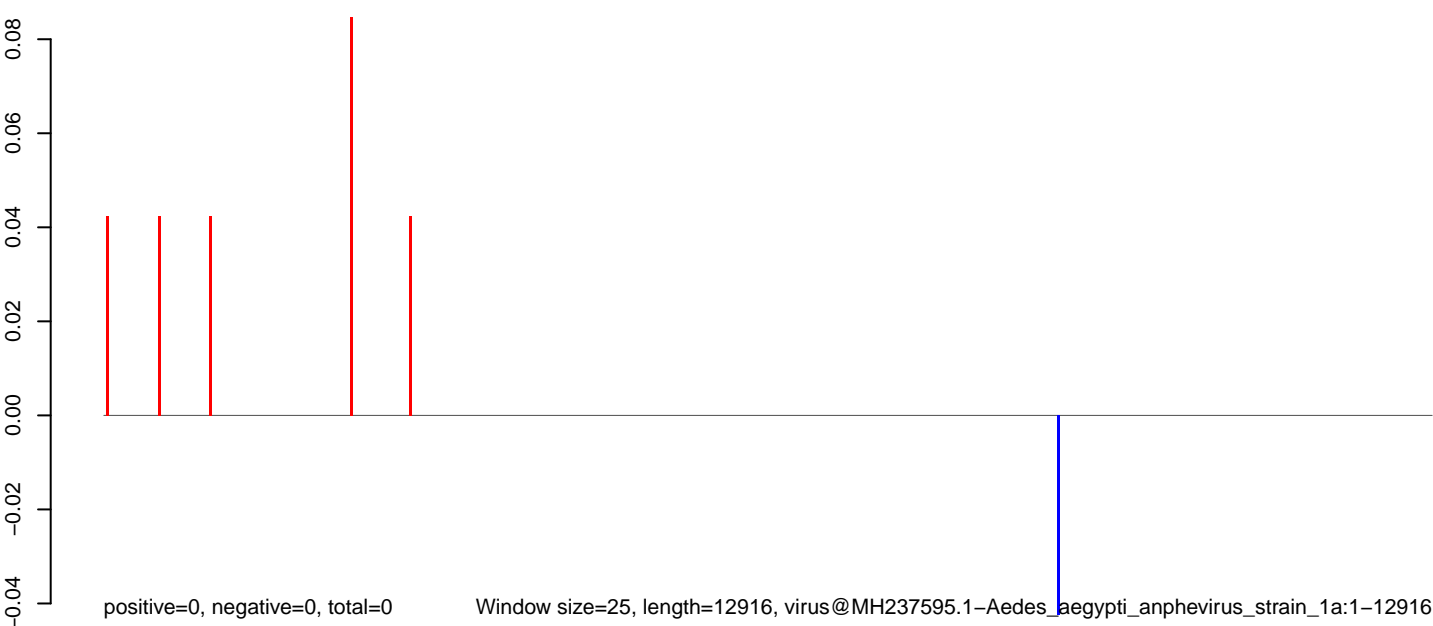
AeAlbo_C636_cells.18_23.rep



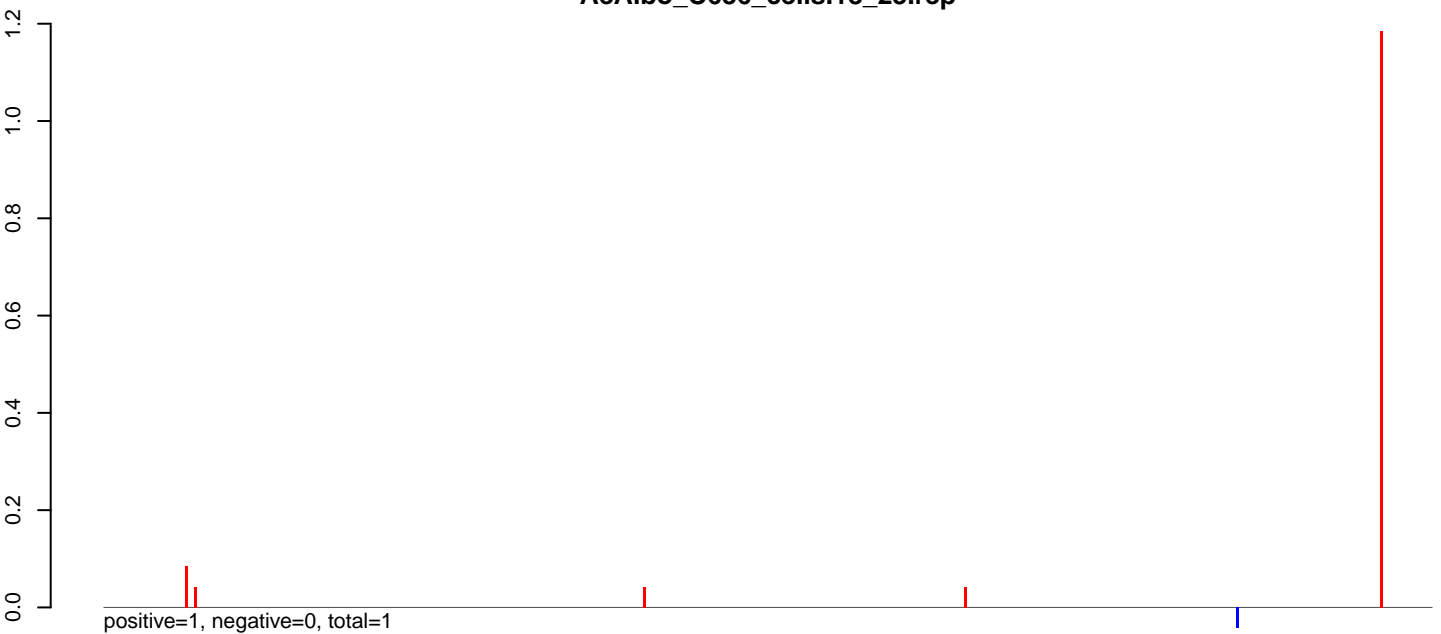
AeAlbo_C636_cells.24_35.rep



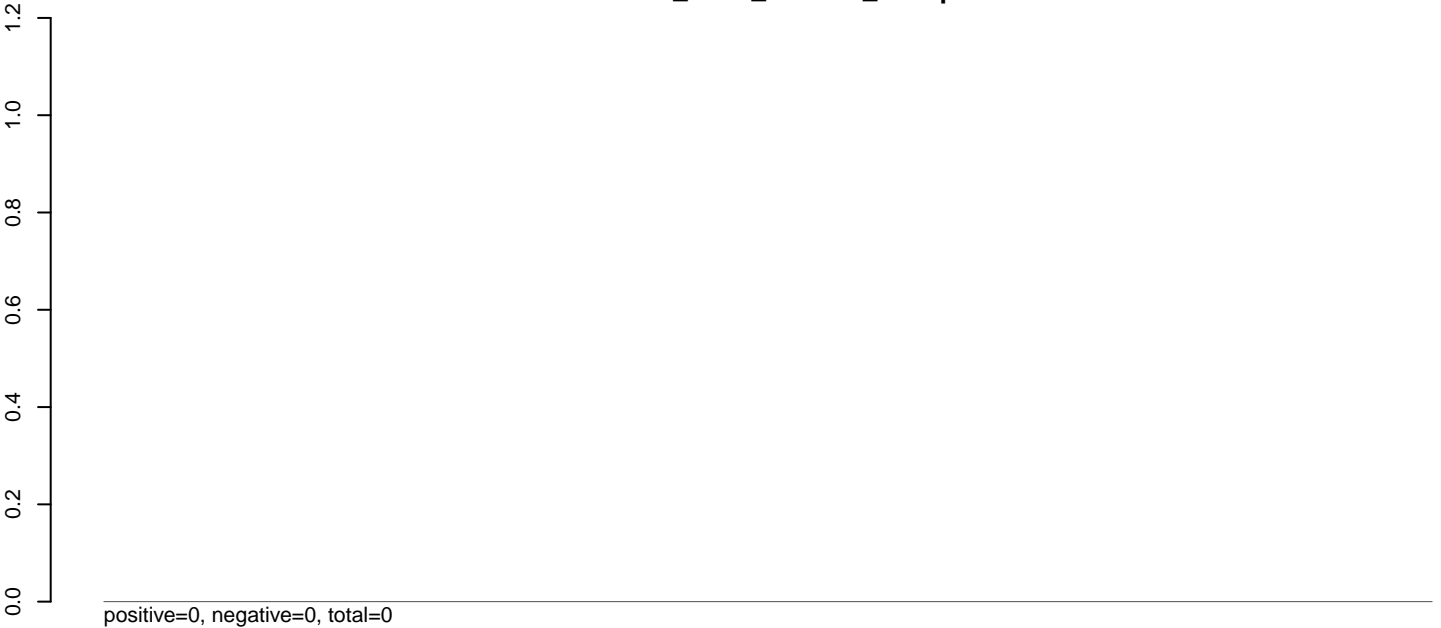
AeAlbo_C636_cells.rep



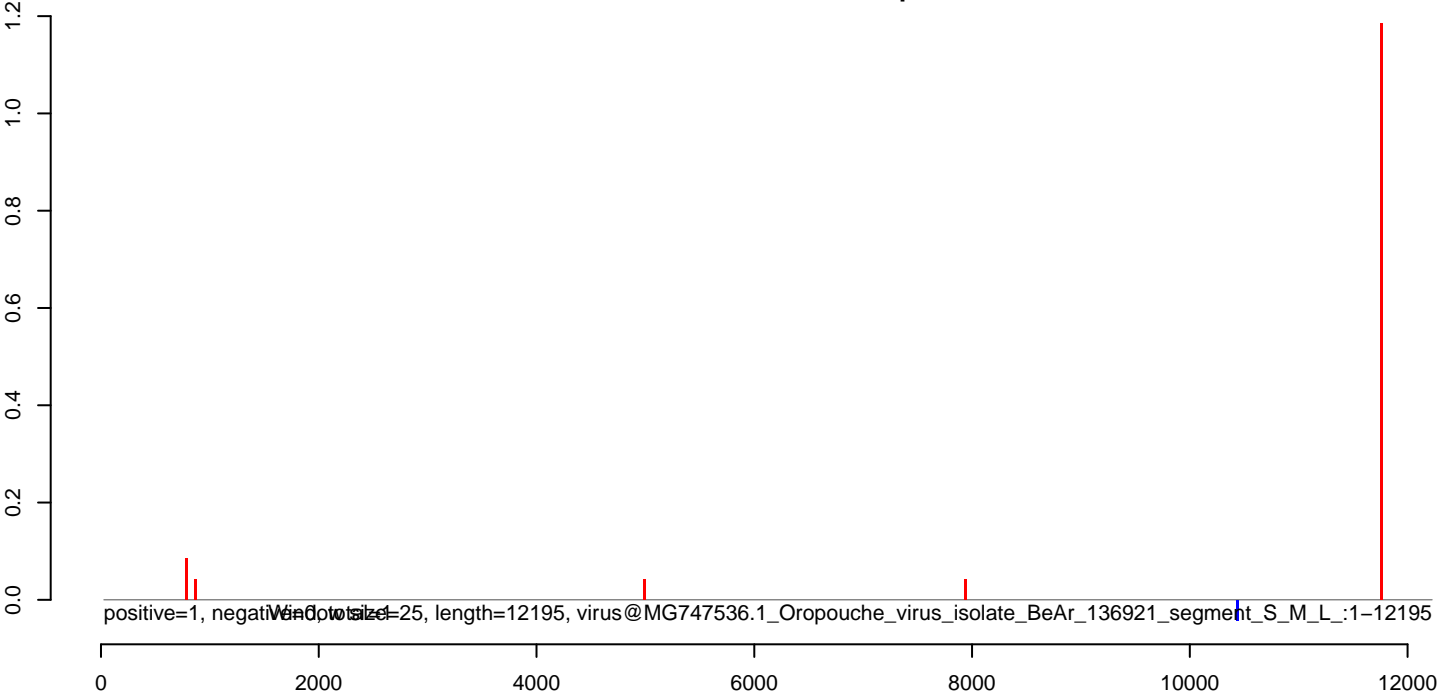
AeAlbo_C636_cells.18_23.rep



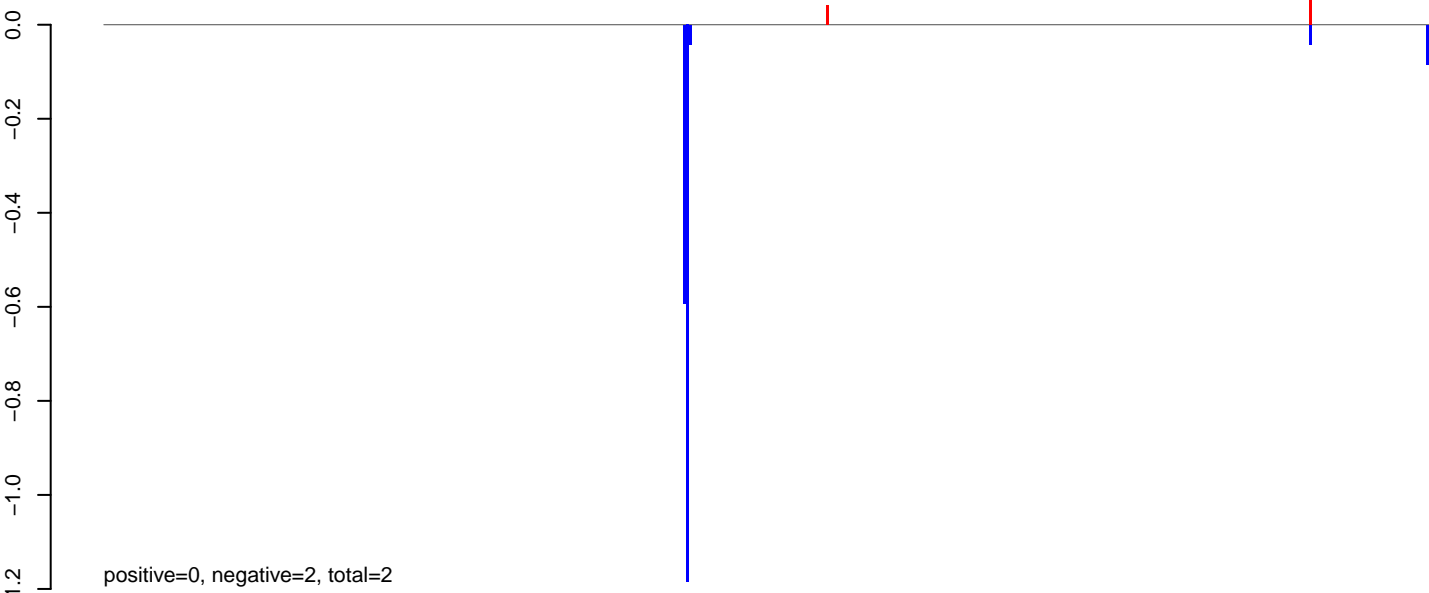
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



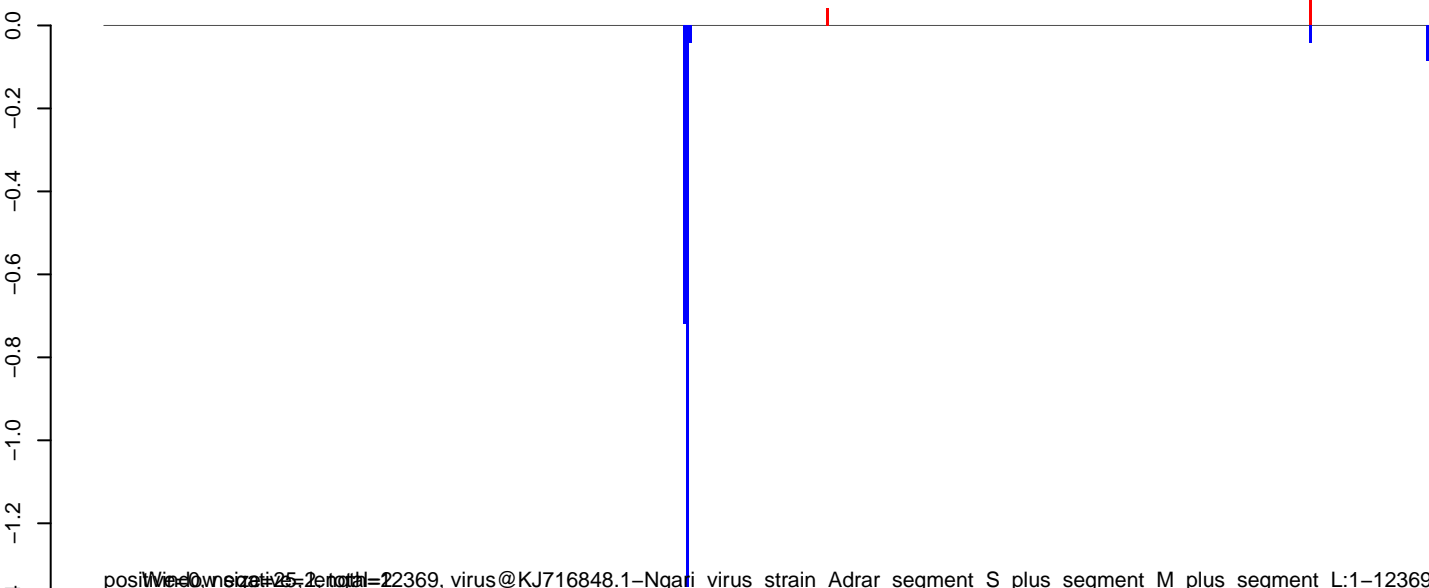
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



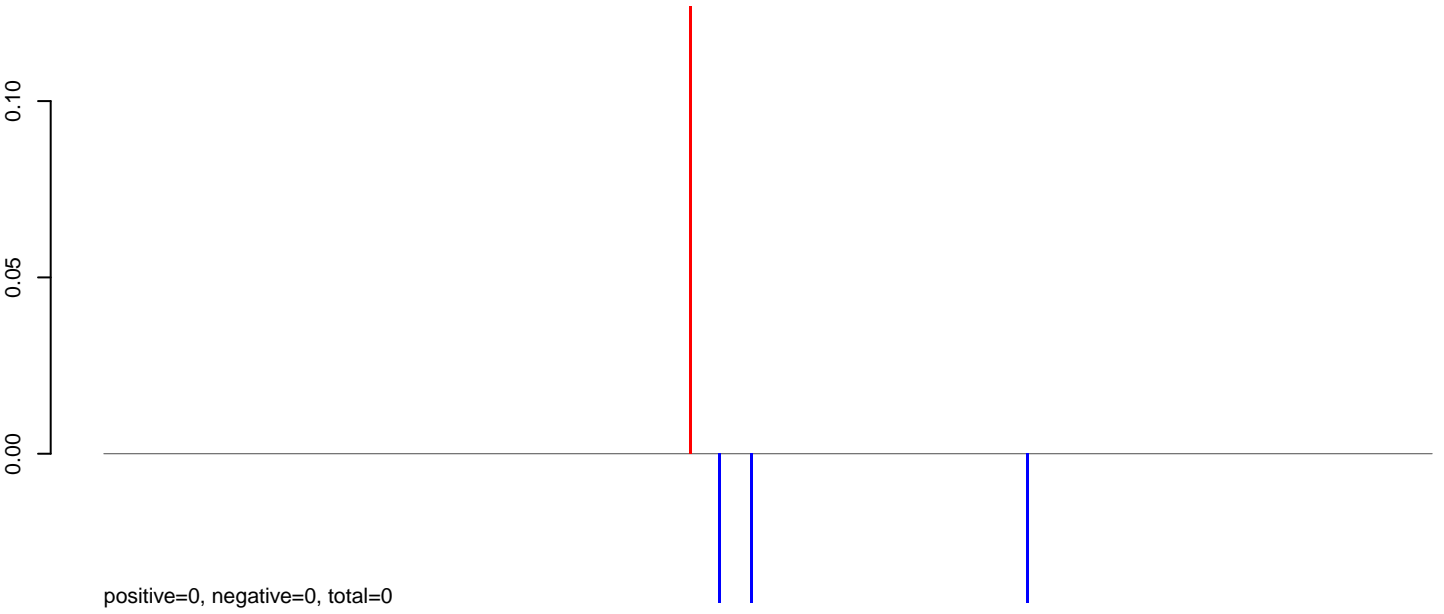
AeAlbo_C636_cells.rep



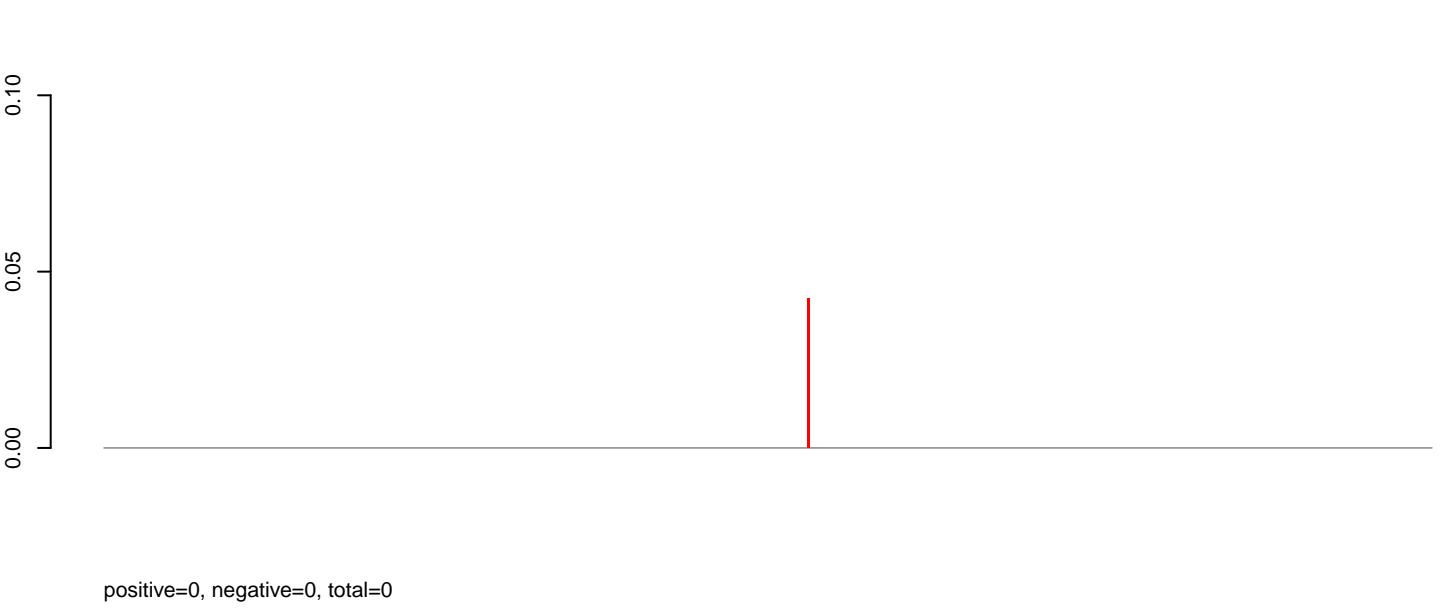
W=0, size=25, length=22369, virus@KJ716848.1-Ngari_virus_strain_Adrar_segment_S_plus_segment_M_plus_segment_L:1-12369

0 2000 4000 6000 8000 10000 12000

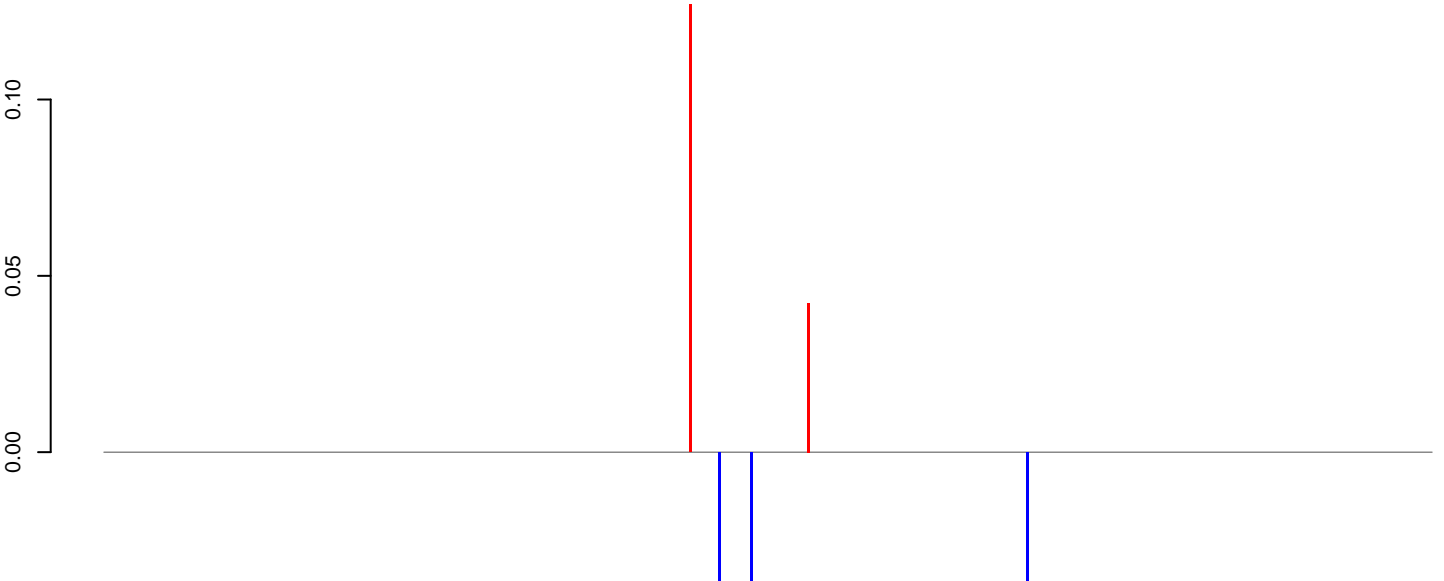
AeAlbo_C636_cells.18_23.rep



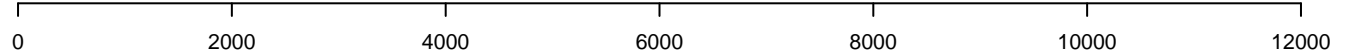
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



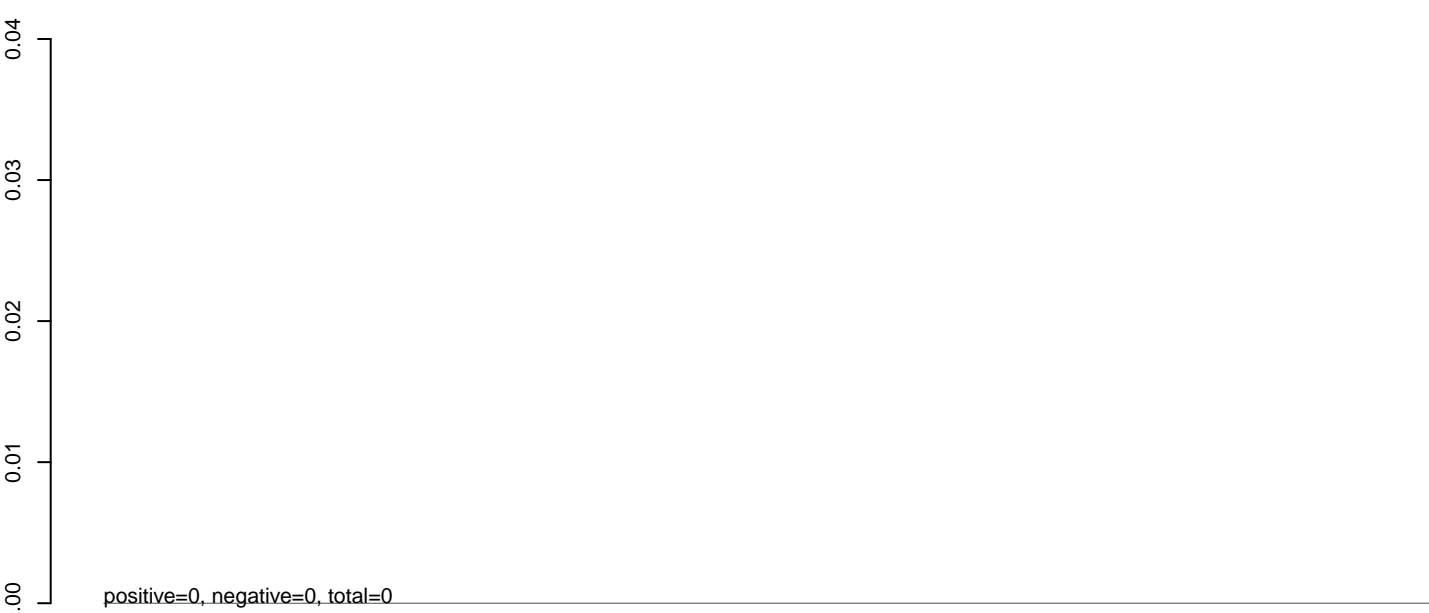
v size=25, l=0, h=400, u=0, c=10
034541.1-Gray_Lodge_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopr:1-12403



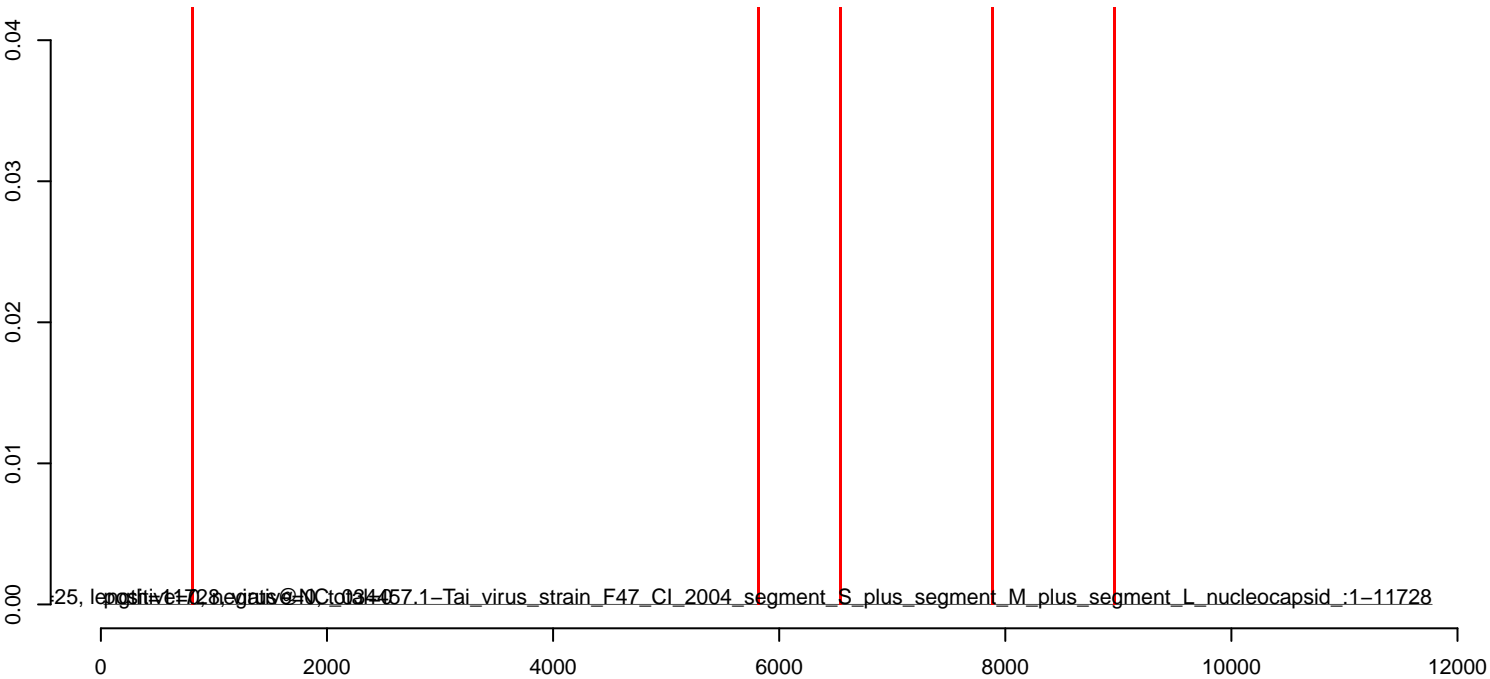
AeAlbo_C636_cells.18_23.rep



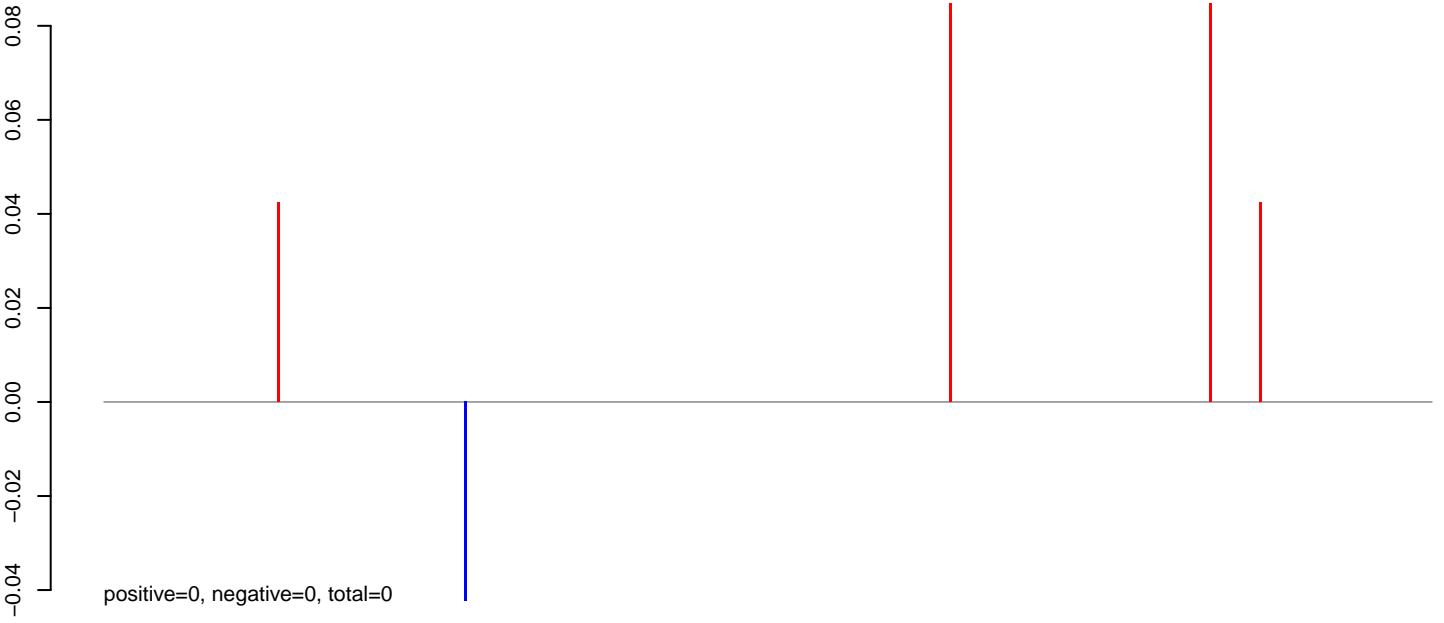
AeAlbo_C636_cells.24_35.rep



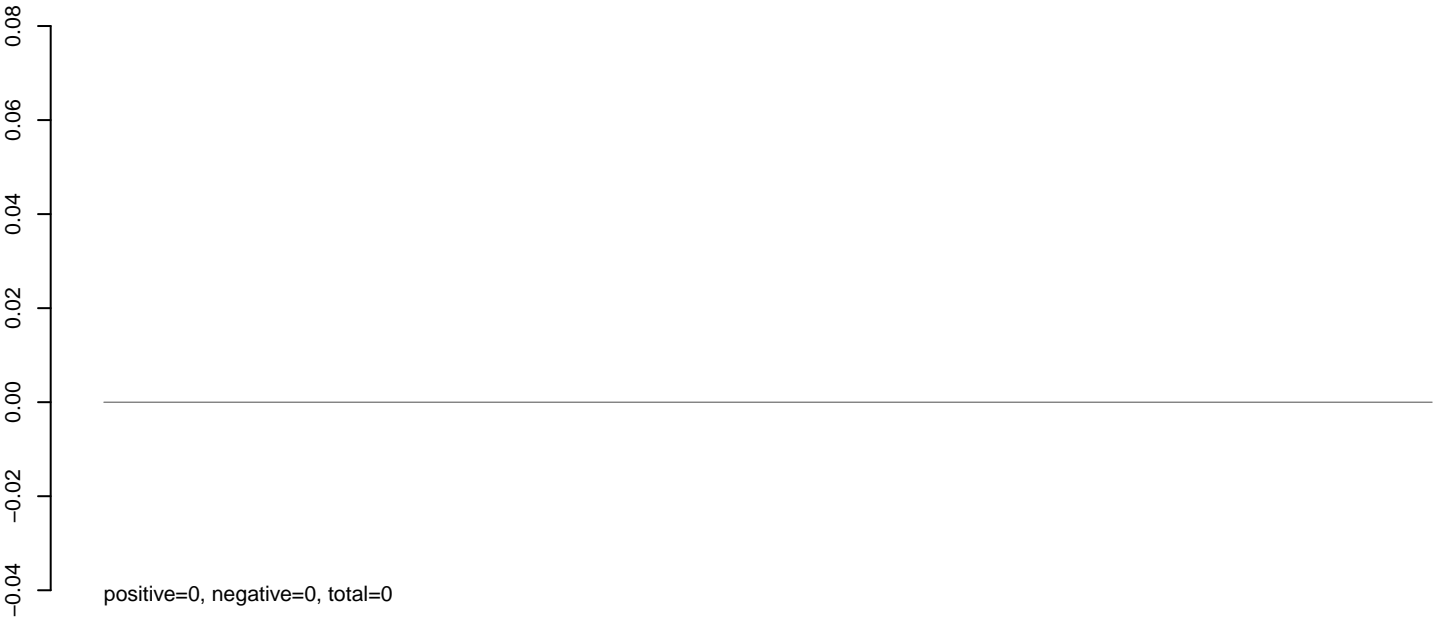
AeAlbo_C636_cells.rep



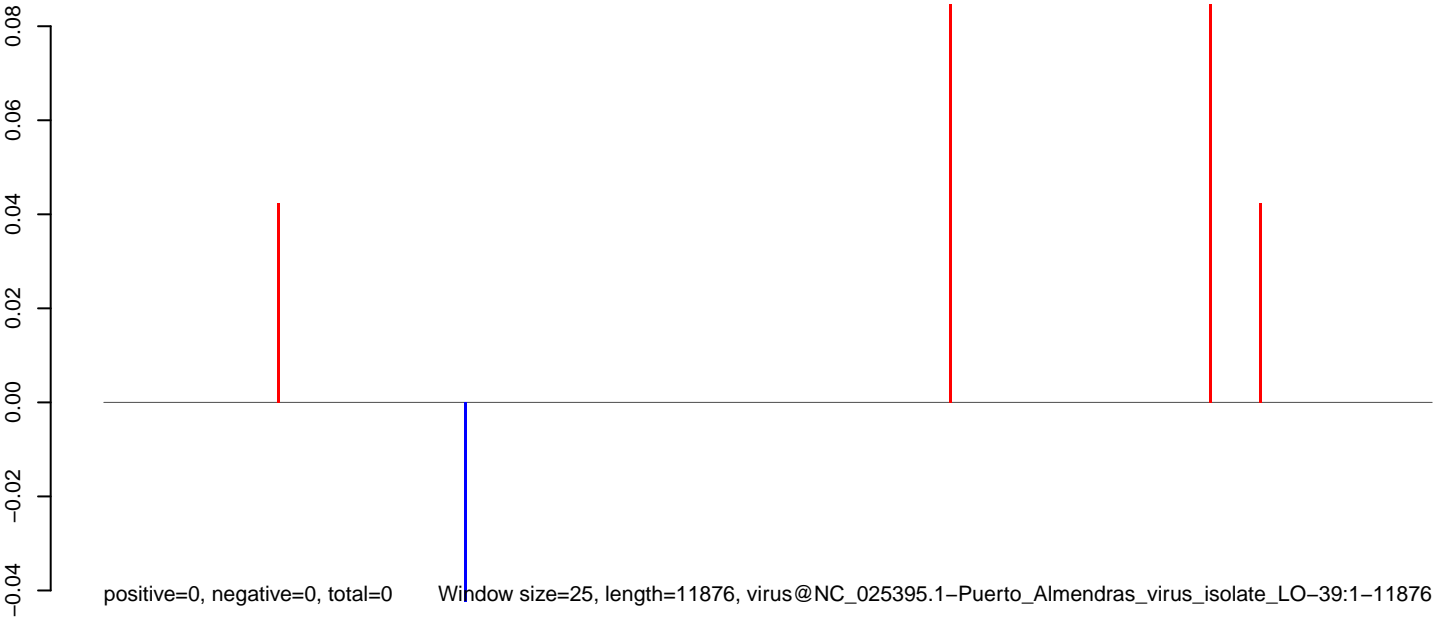
AeAlbo_C636_cells.18_23.rep



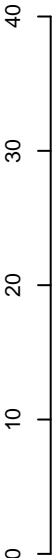
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

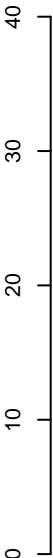


AeAlbo_C636_cells.18_23.rep



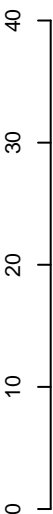
positive=118, negative=6, total=124

AeAlbo_C636_cells.24_35.rep

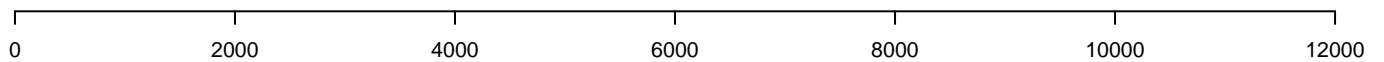


positive=16, negative=0, total=16

AeAlbo_C636_cells.rep



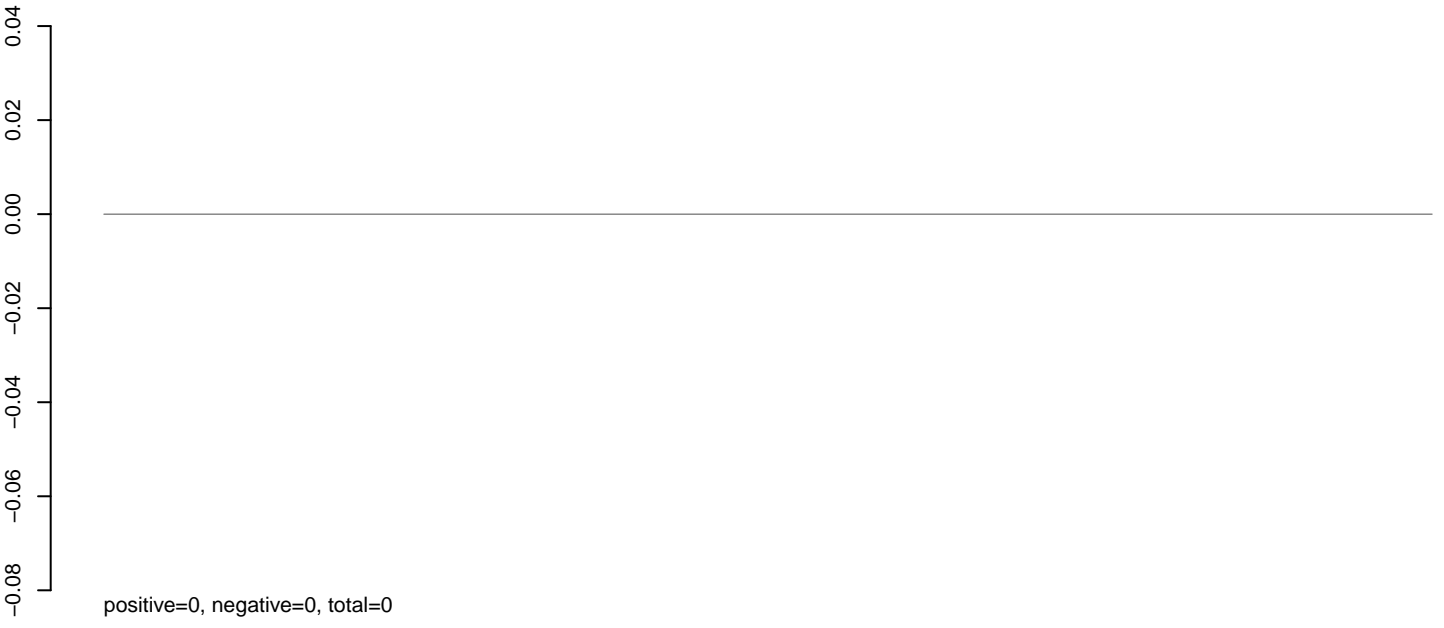
downloaded from https://www.ncbi.nlm.nih.gov/trace/1147991.1-Trocaravirus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotein:1-12052



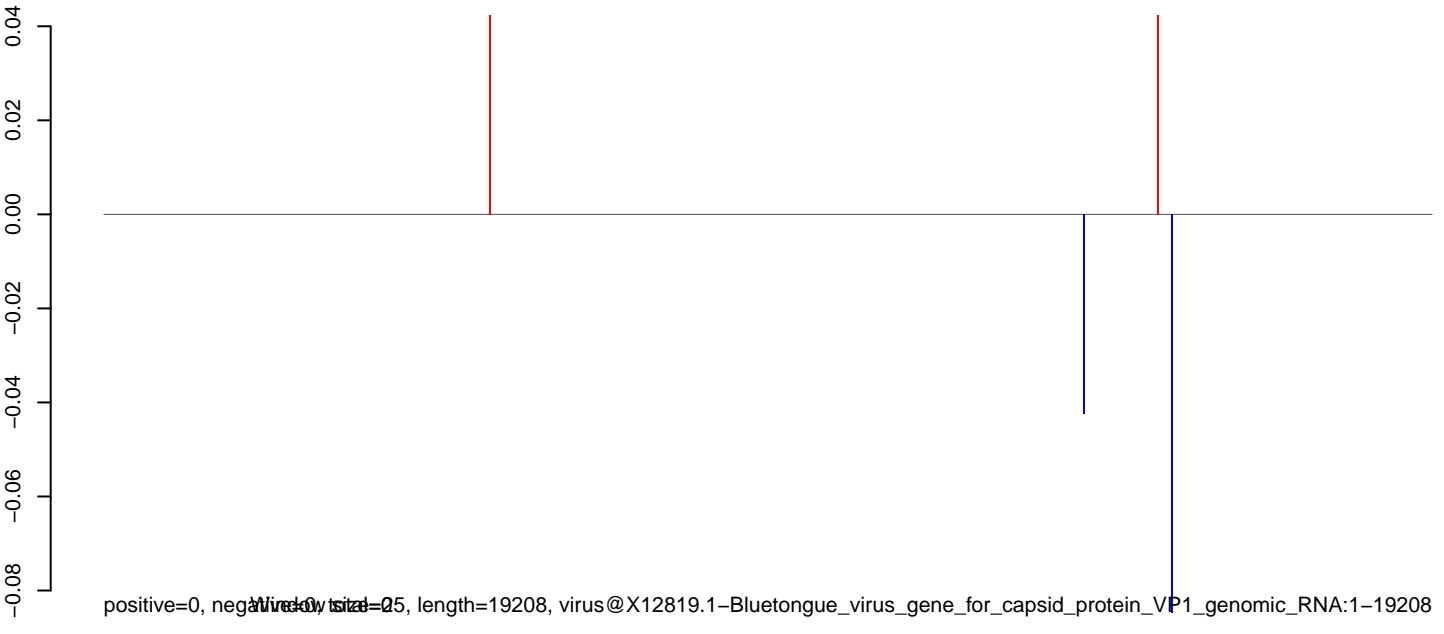
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



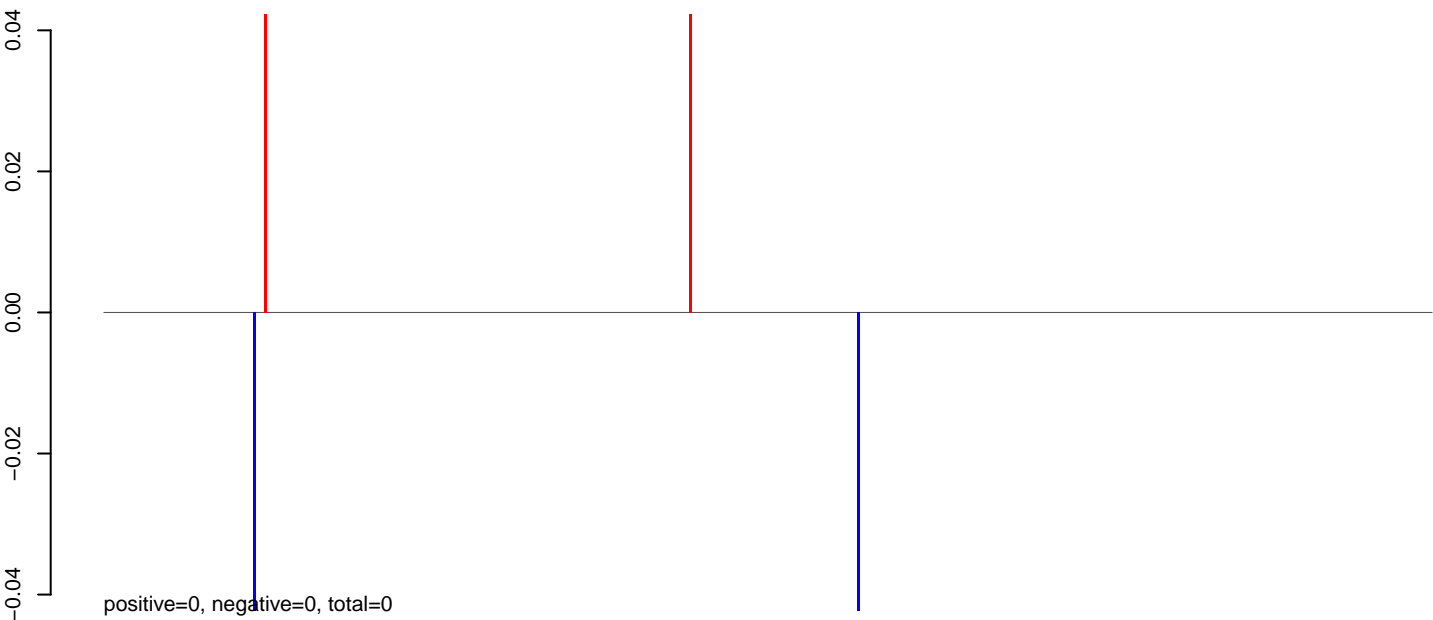
AeAlbo_C636_cells.rep



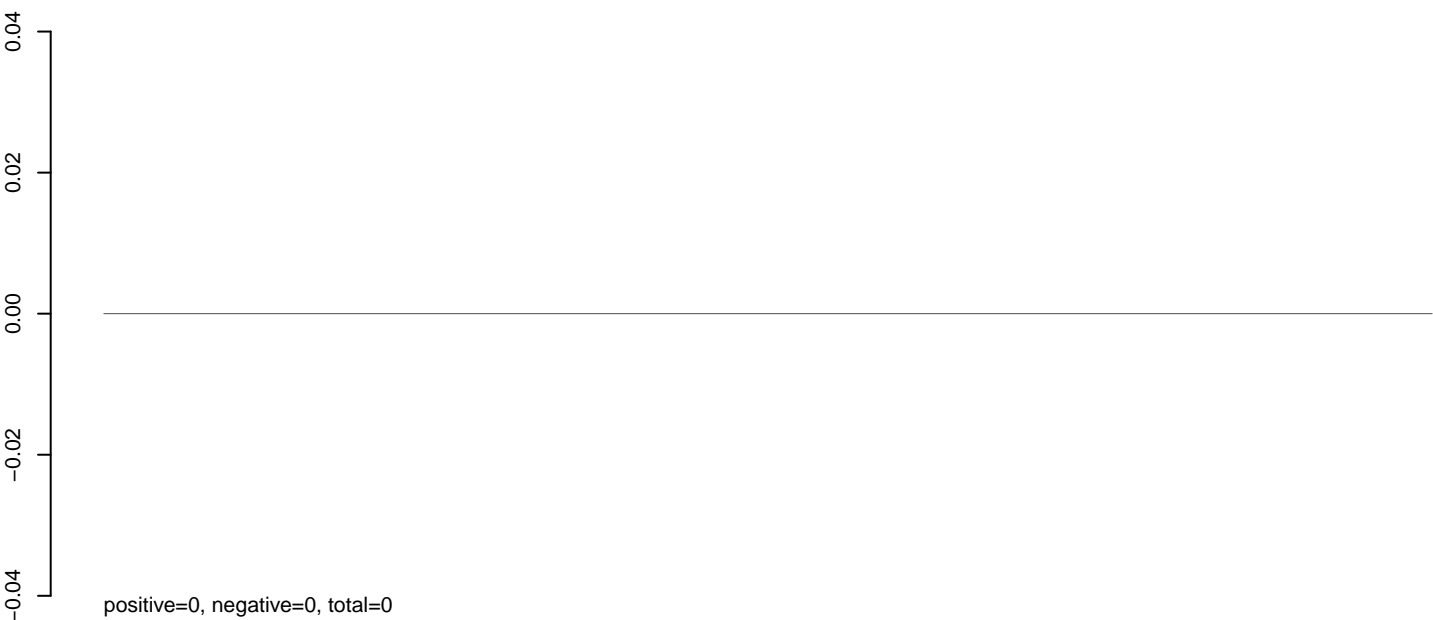
virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208

0 5000 10000 15000 20000

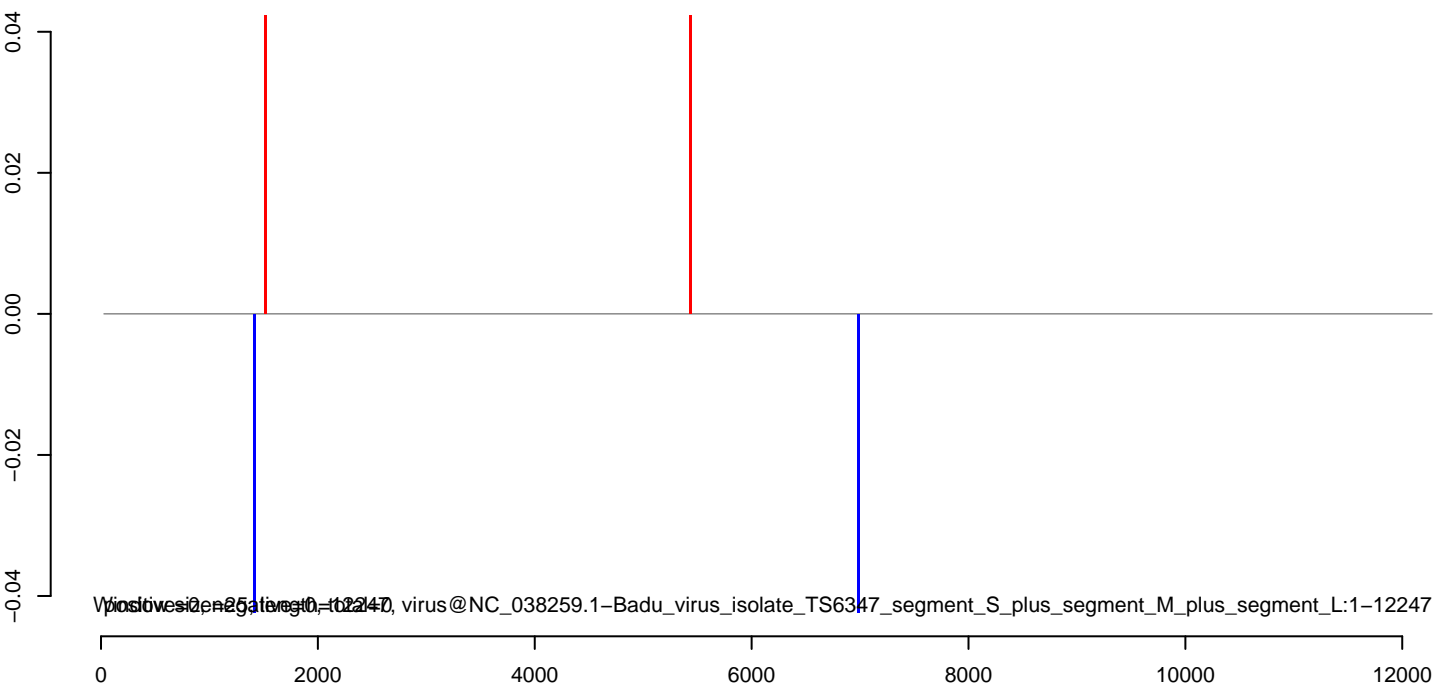
AeAlbo_C636_cells.18_23.rep



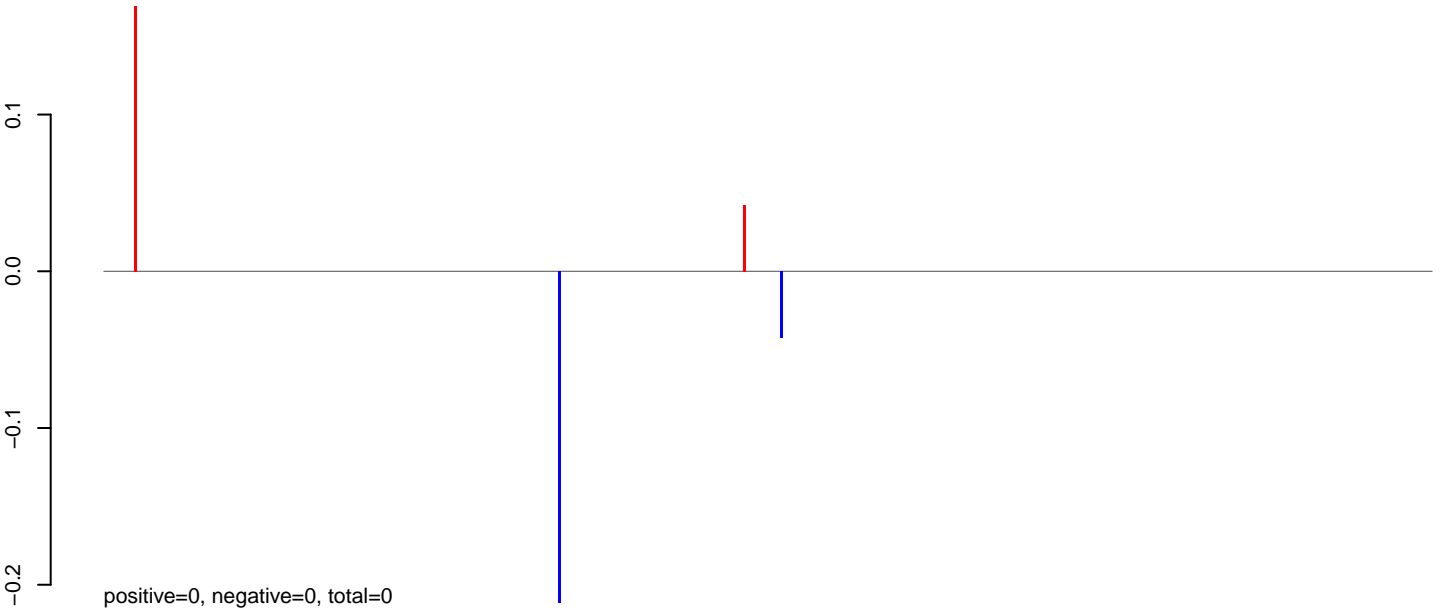
AeAlbo_C636_cells.24_35.rep



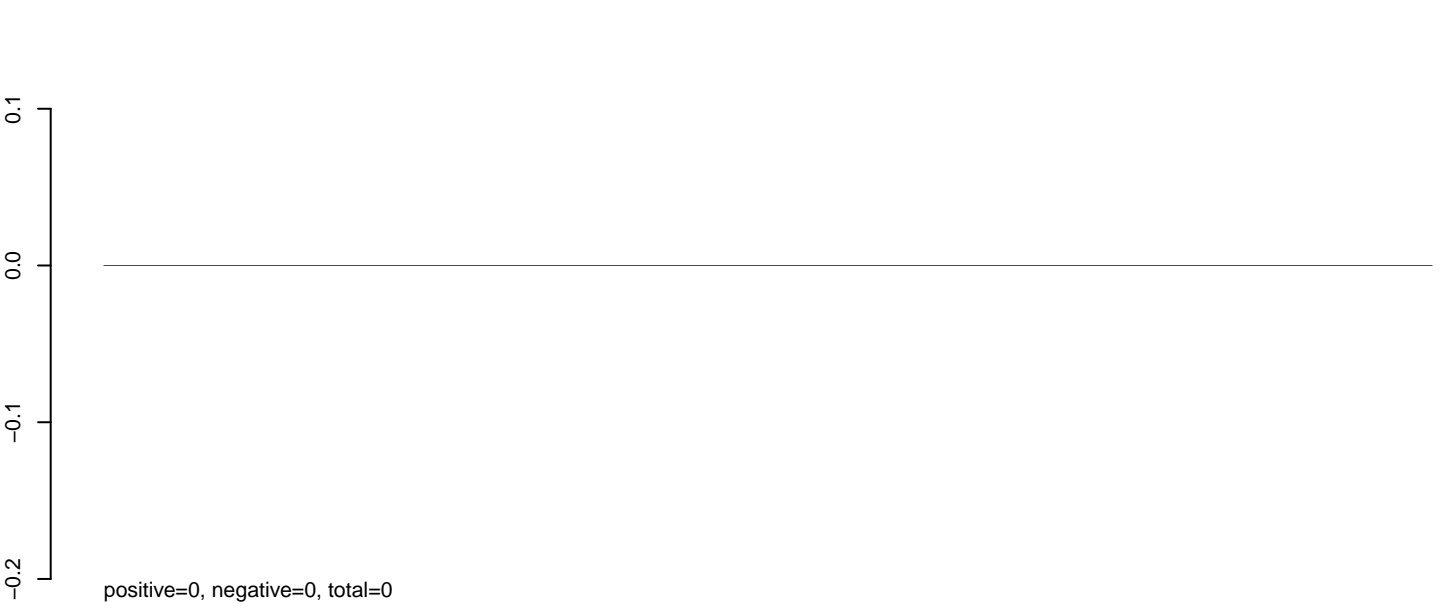
AeAlbo_C636_cells.rep



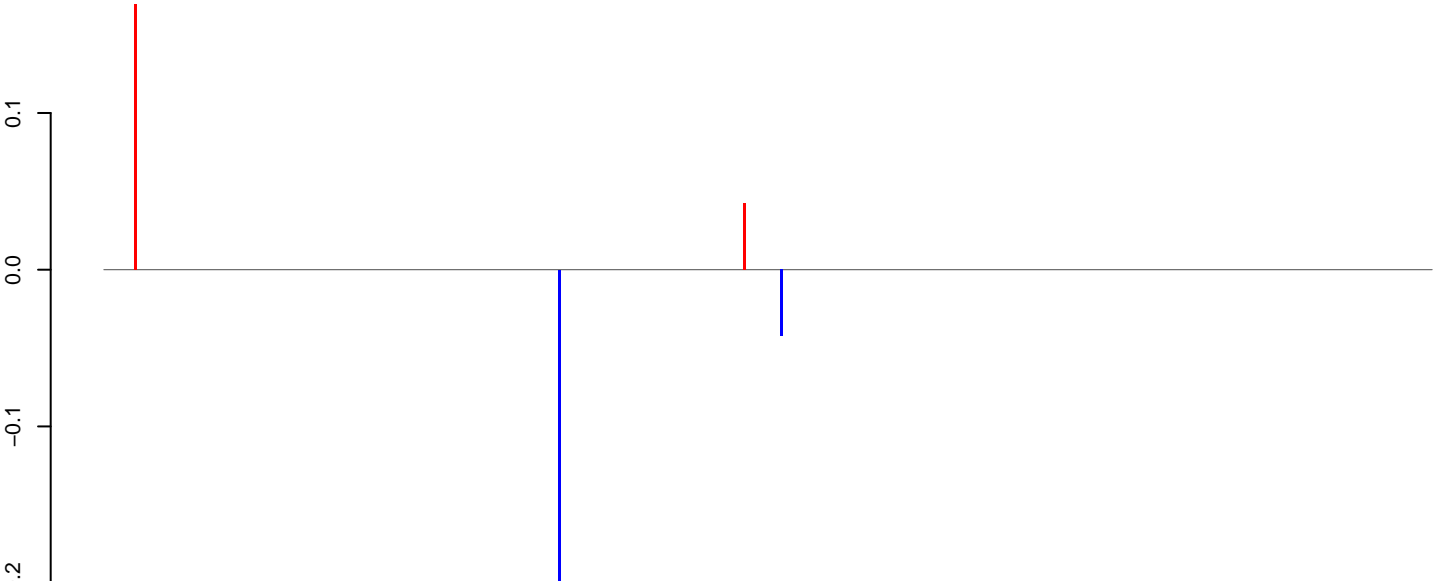
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



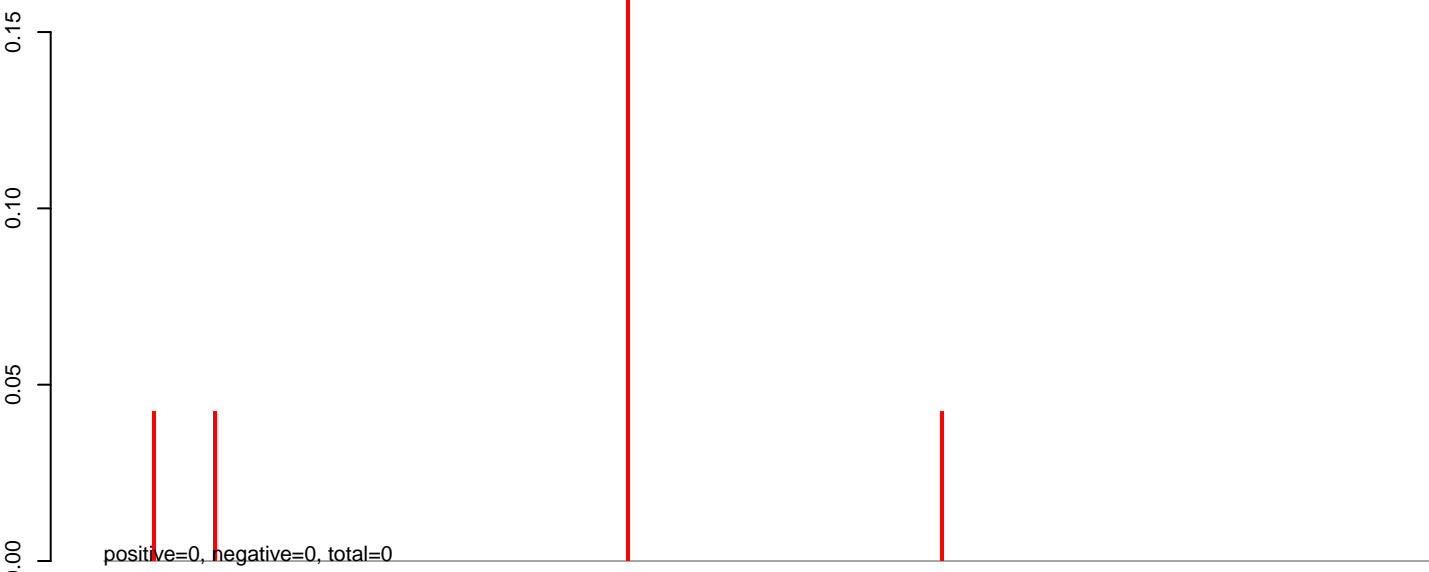
AeAlbo_C636_cells.rep



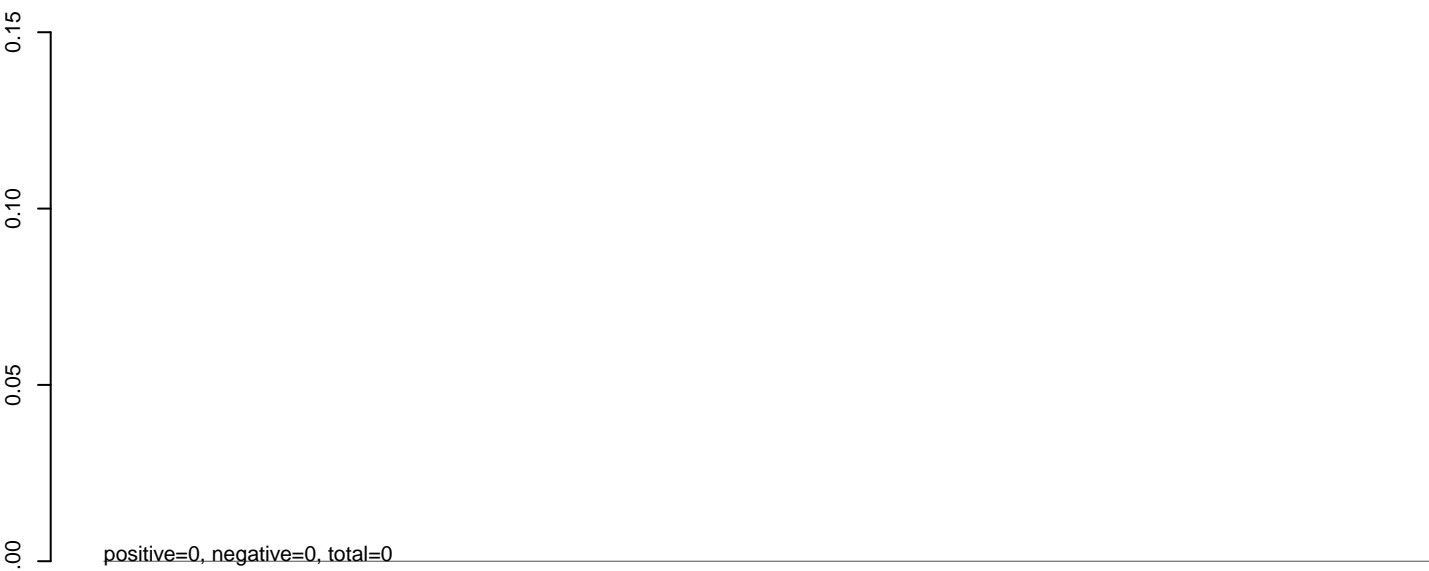
bow size 25, length 1294, virus NC_034530.1-Marco_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_proteins:1-13294

0 2000 4000 6000 8000 10000 12000

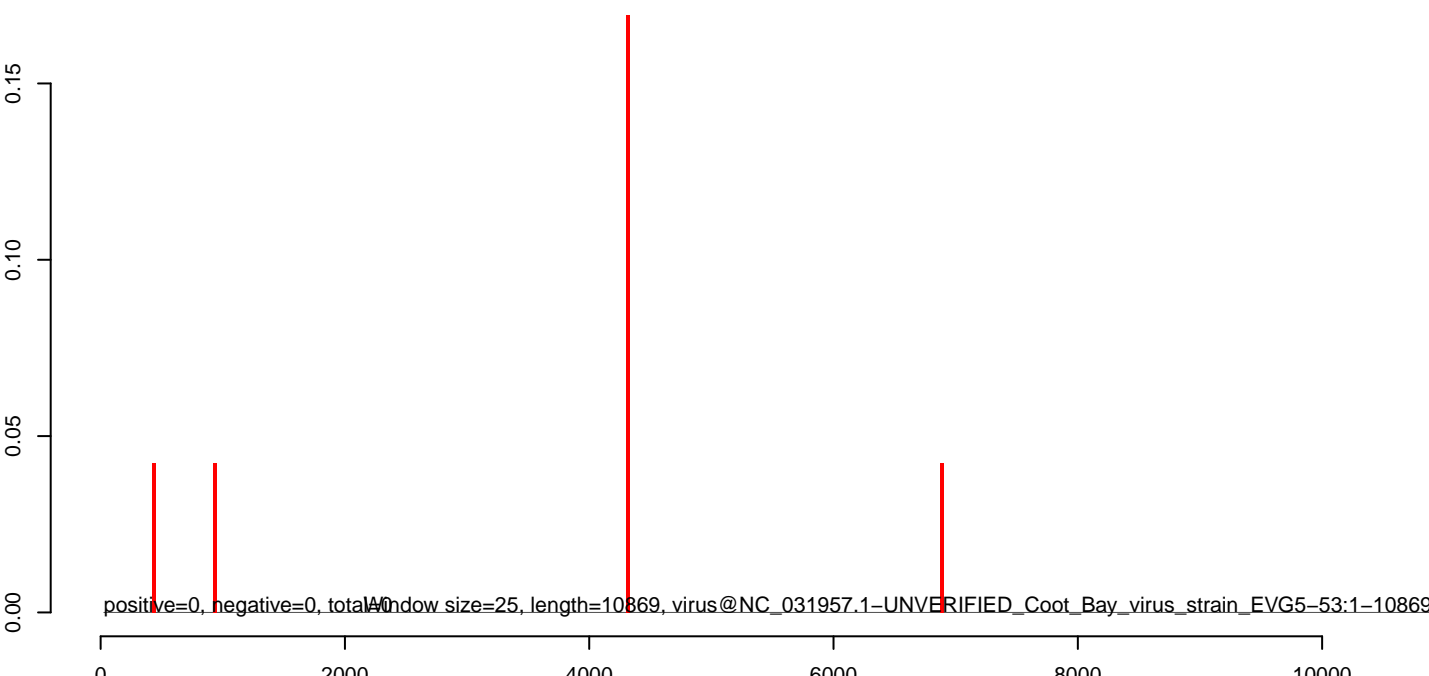
AeAlbo_C636_cells.18_23.rep



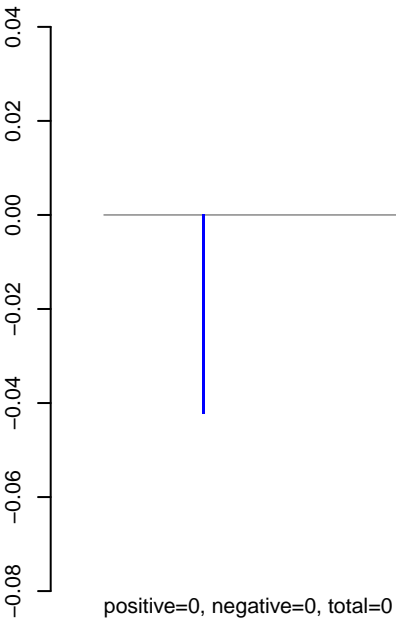
AeAlbo_C636_cells.24_35.rep



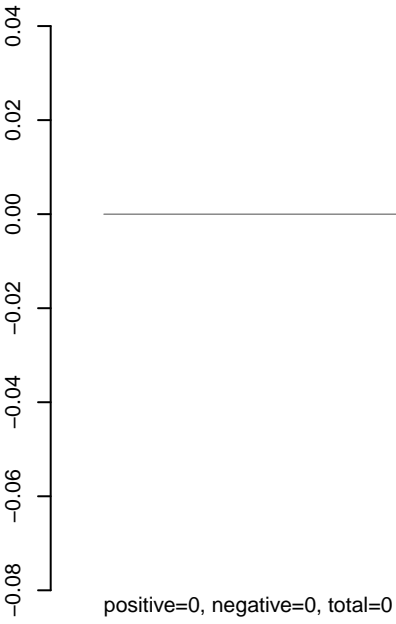
AeAlbo_C636_cells.rep



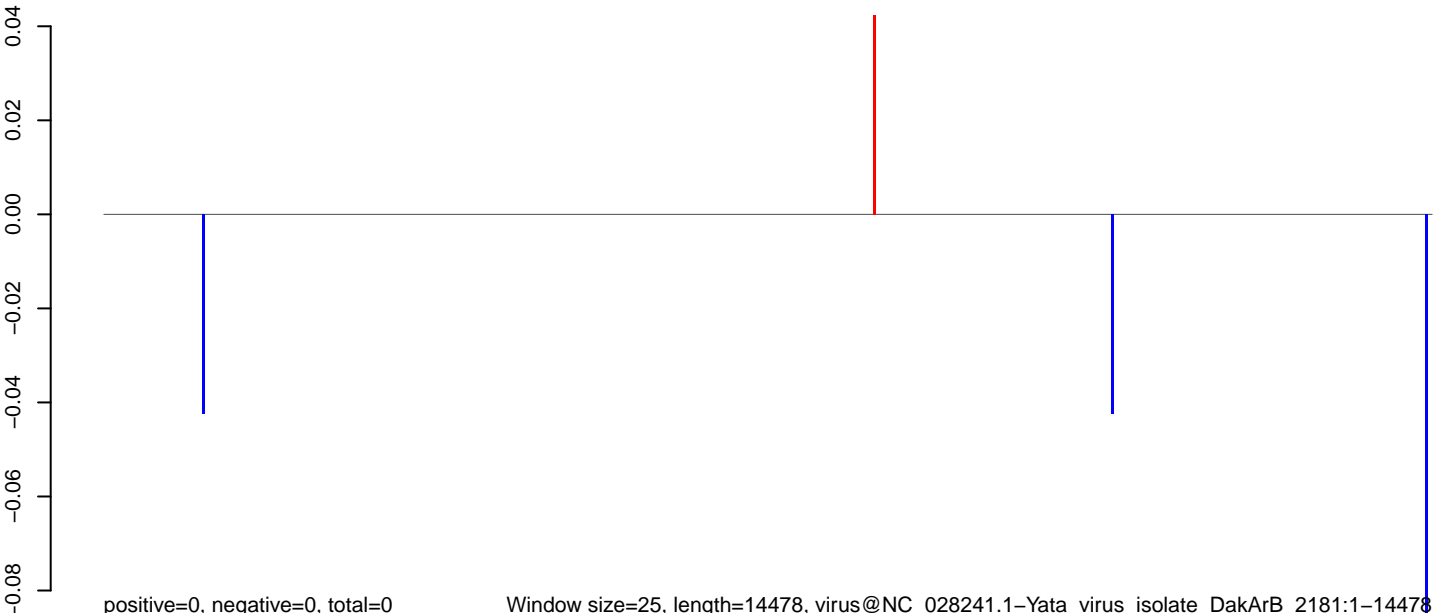
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



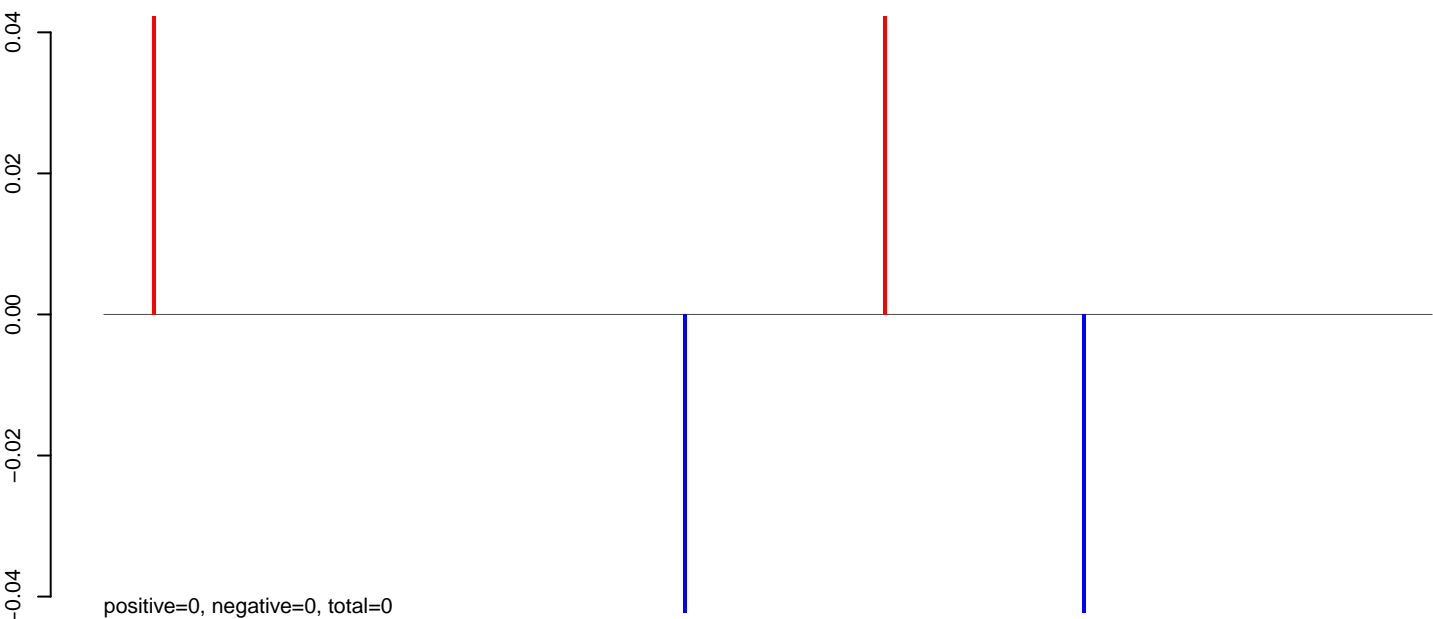
AeAlbo_C636_cells.rep



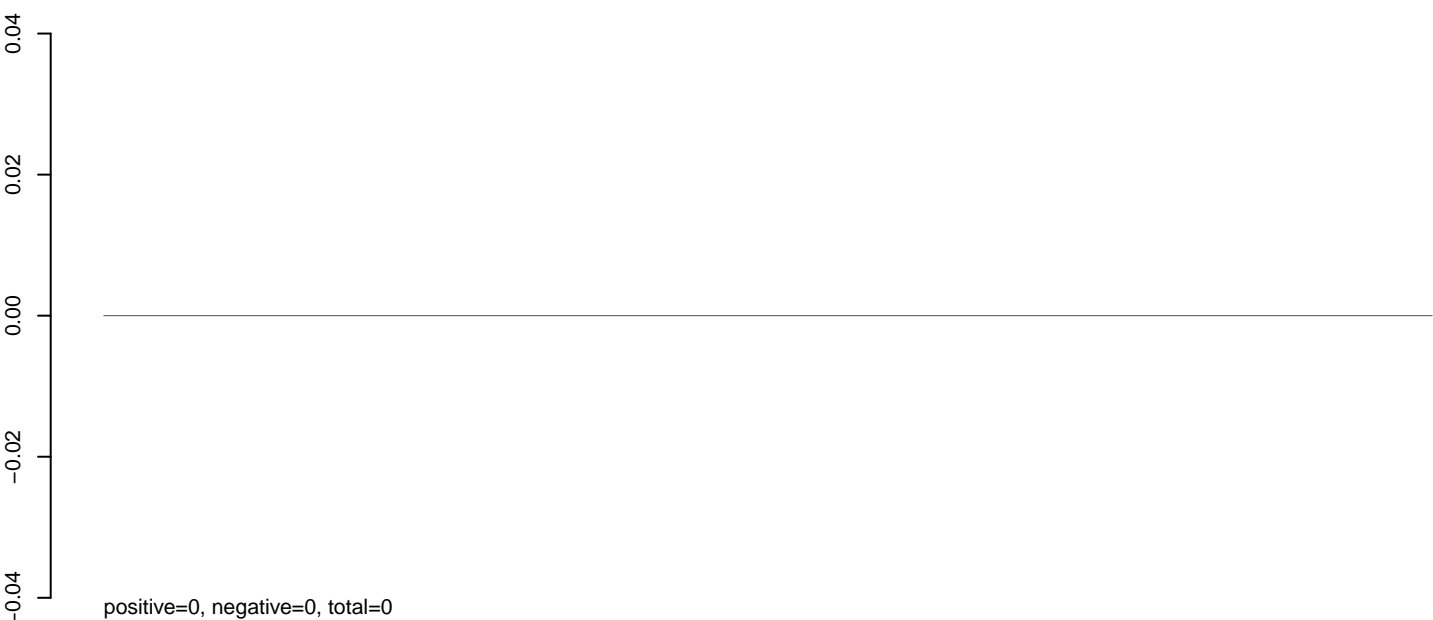
Window size=25, length=14478, virus@NC_028241.1-Yata_virus_isolate_DakArB_2181:1-14478

0 5000 10000 15000

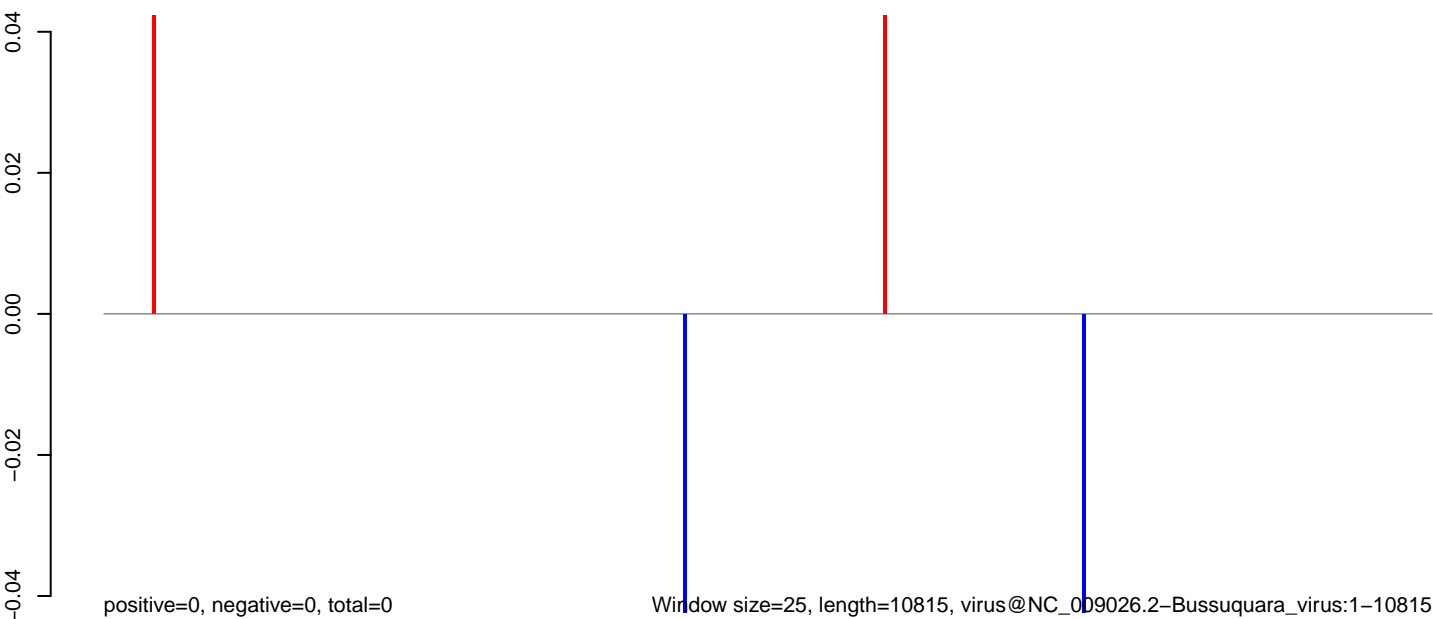
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

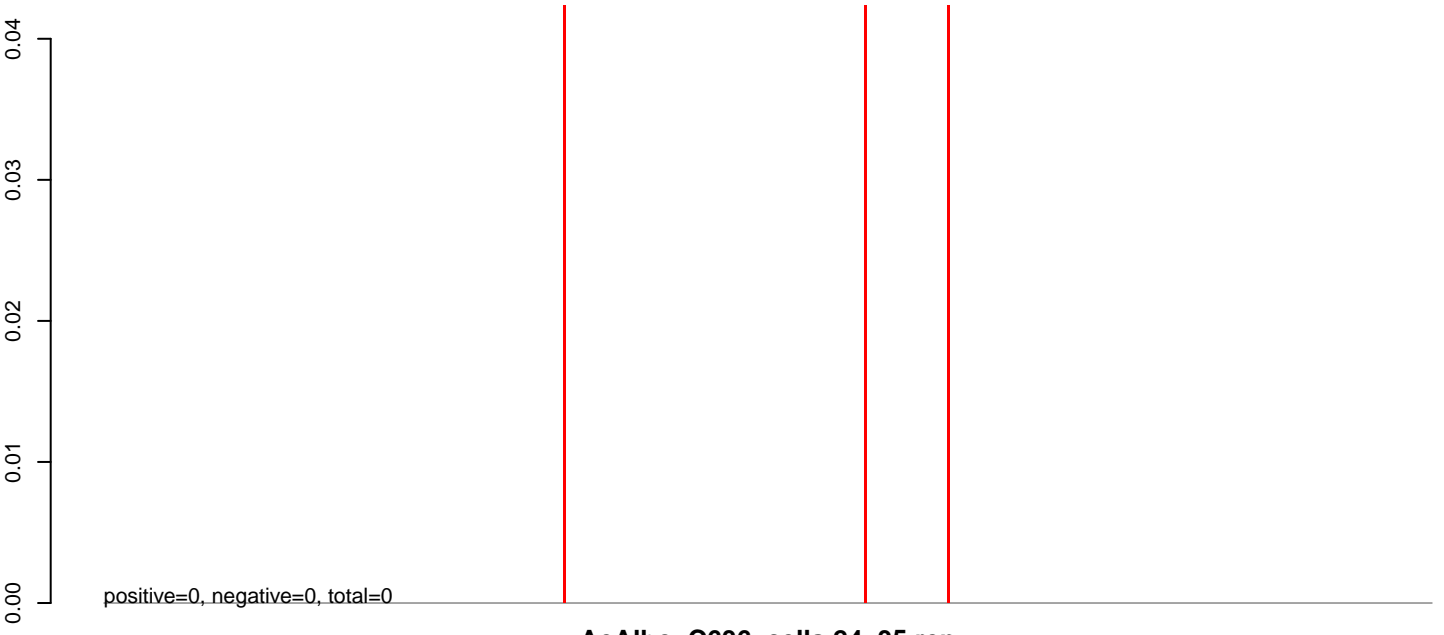


AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

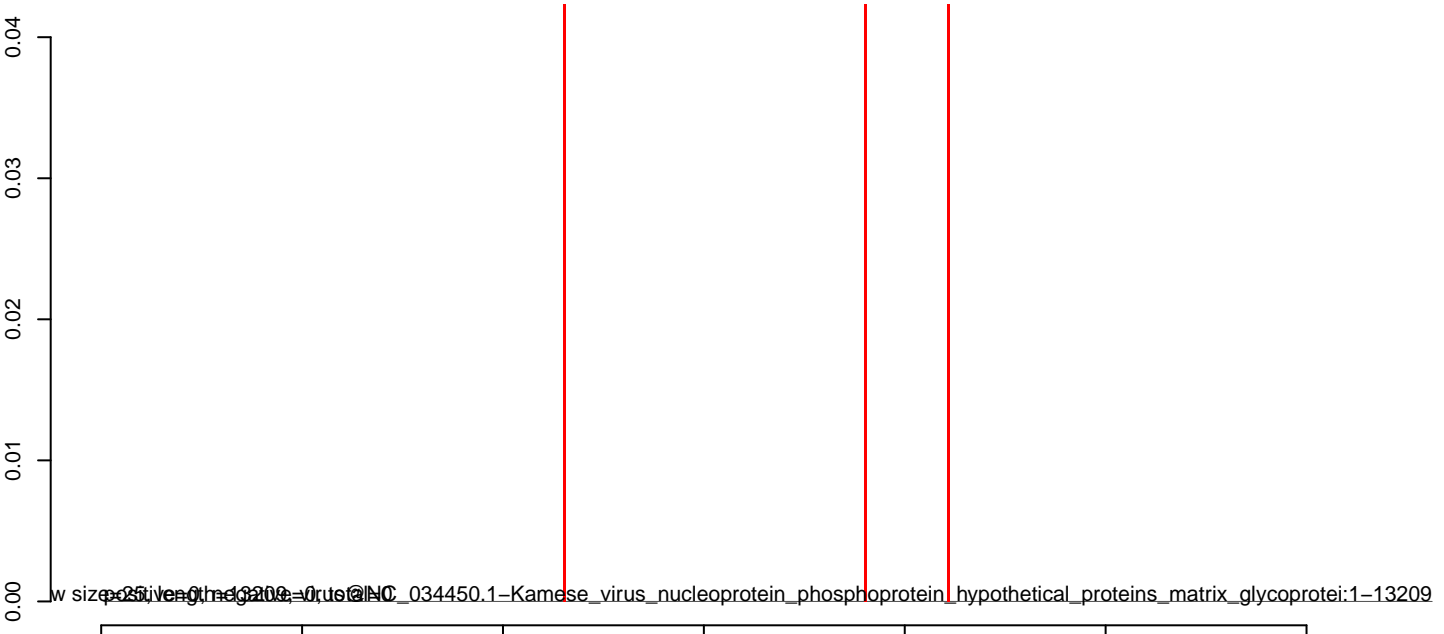
AeAlbo_C636_cells.18_23.rep



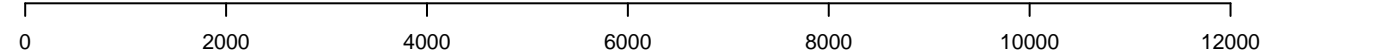
AeAlbo_C636_cells.24_35.rep



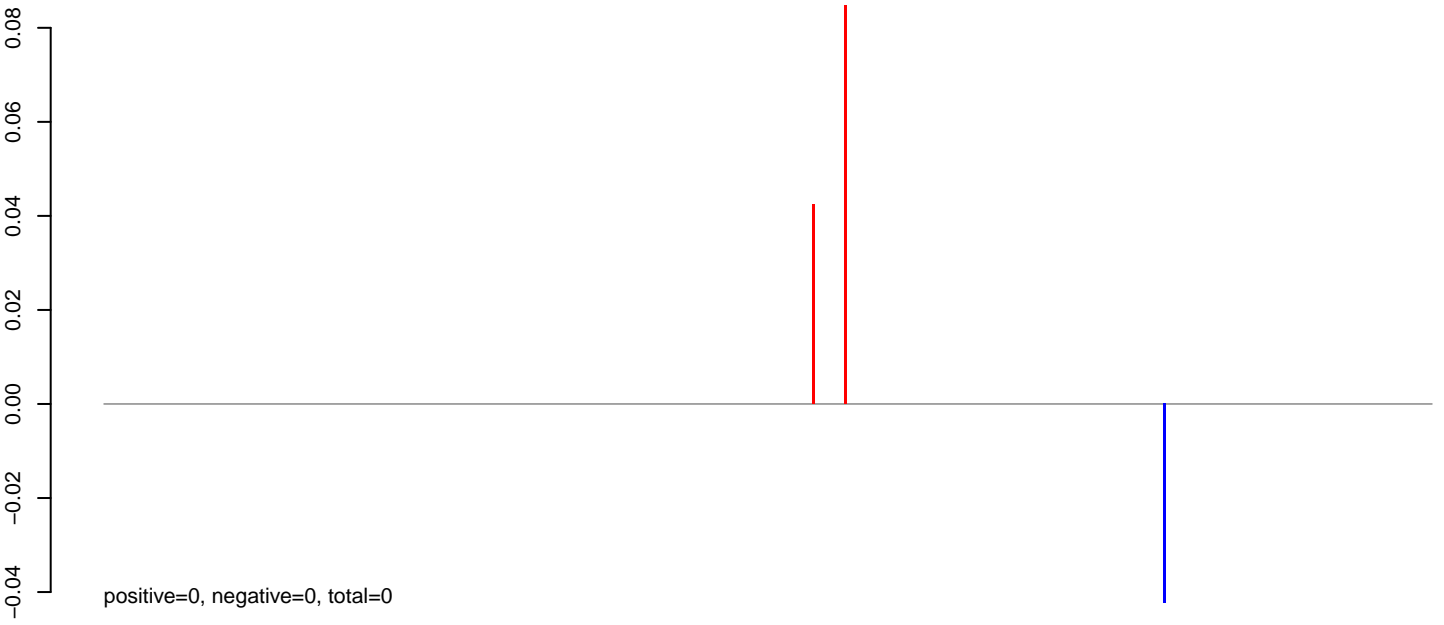
AeAlbo_C636_cells.rep



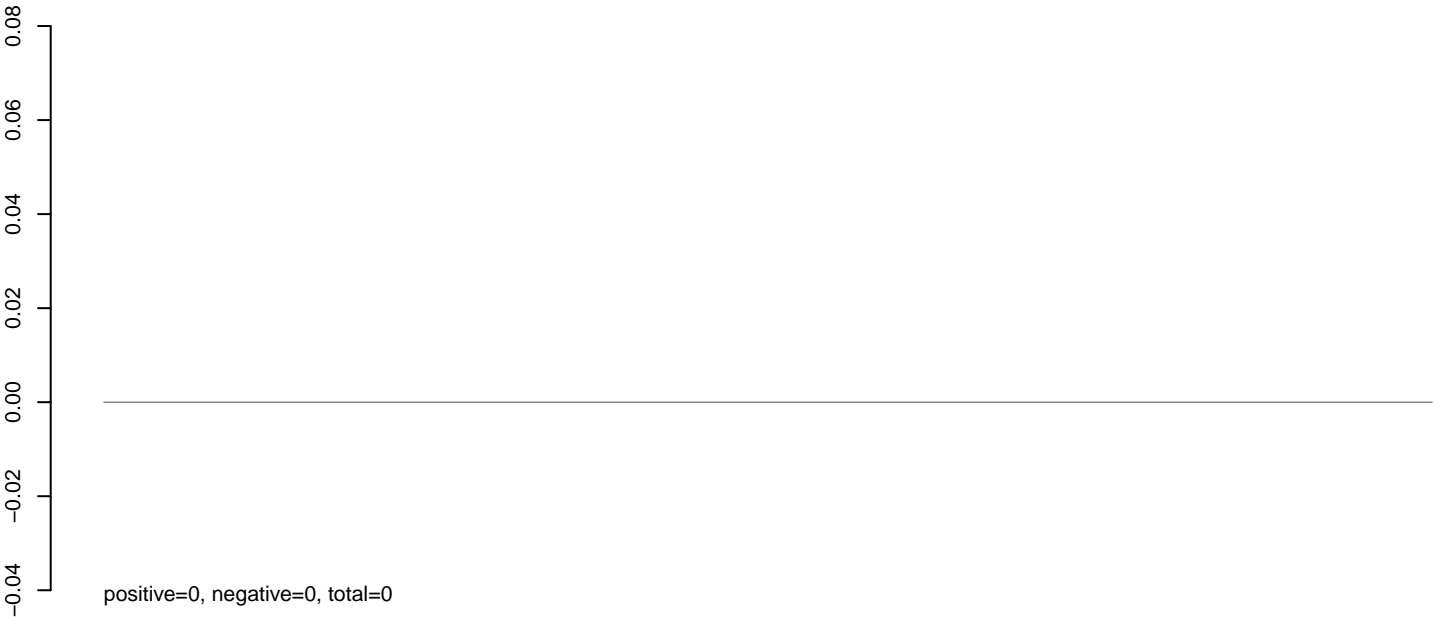
size=25, length=13200, use@NC_034450.1-Kamese_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprotei:1-13209



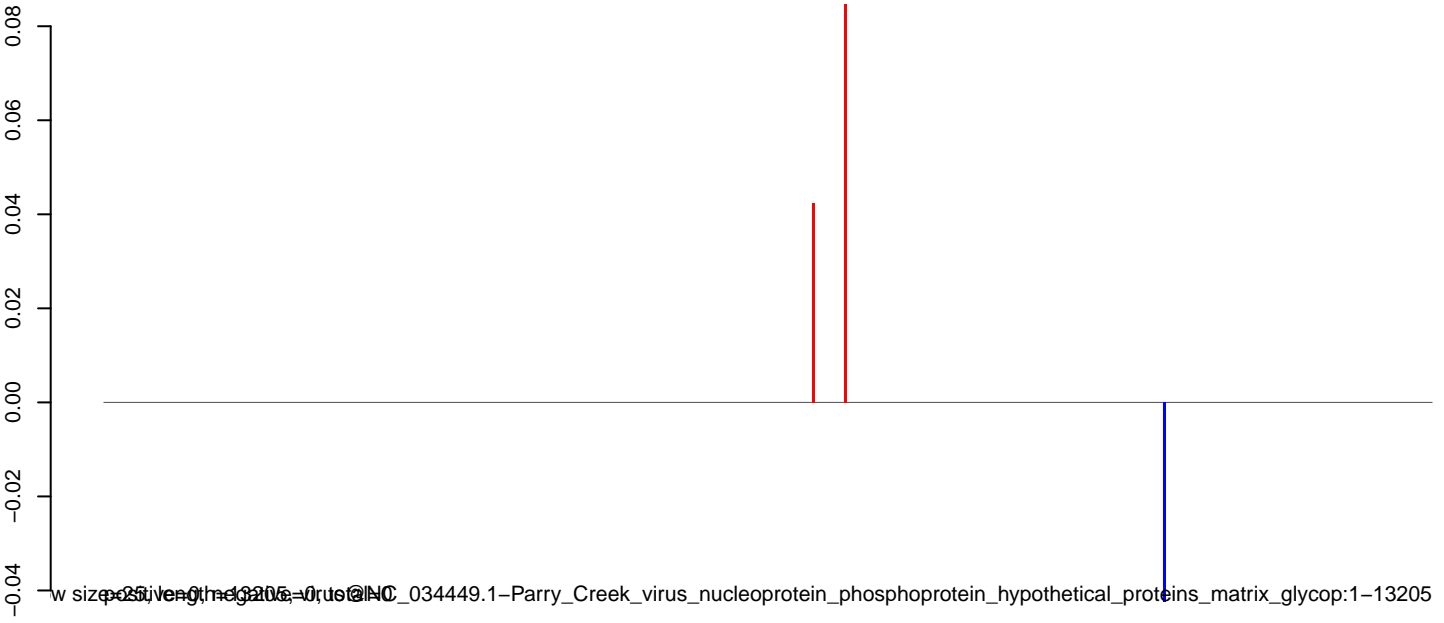
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



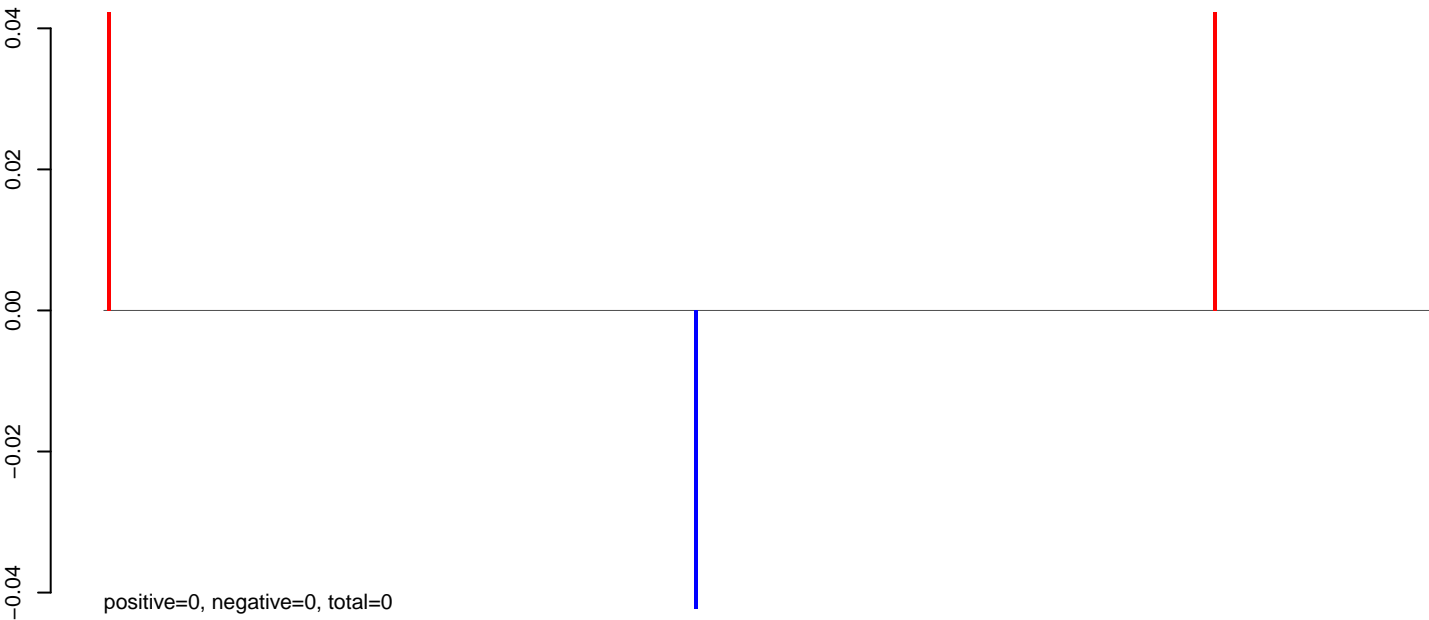
AeAlbo_C636_cells.rep



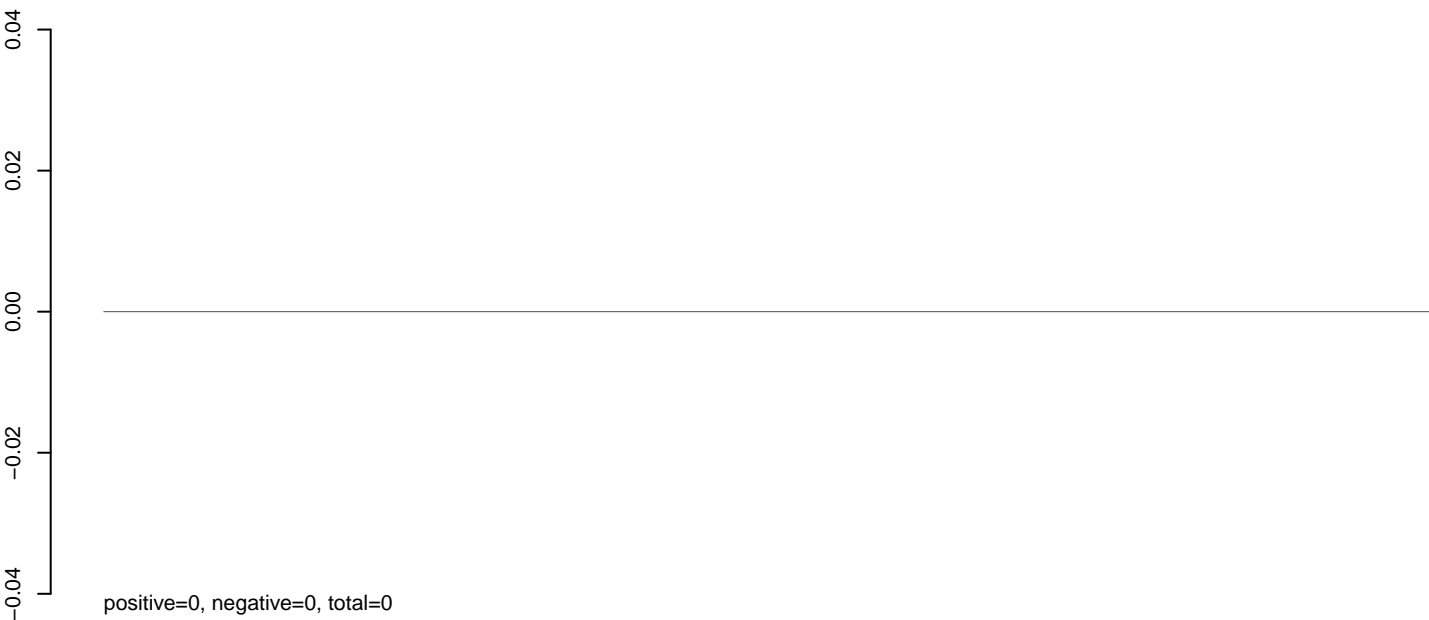
size=25, length=13205, use@NC_034449.1-Parry_Creek_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycop:1-13205

0 2000 4000 6000 8000 10000 12000

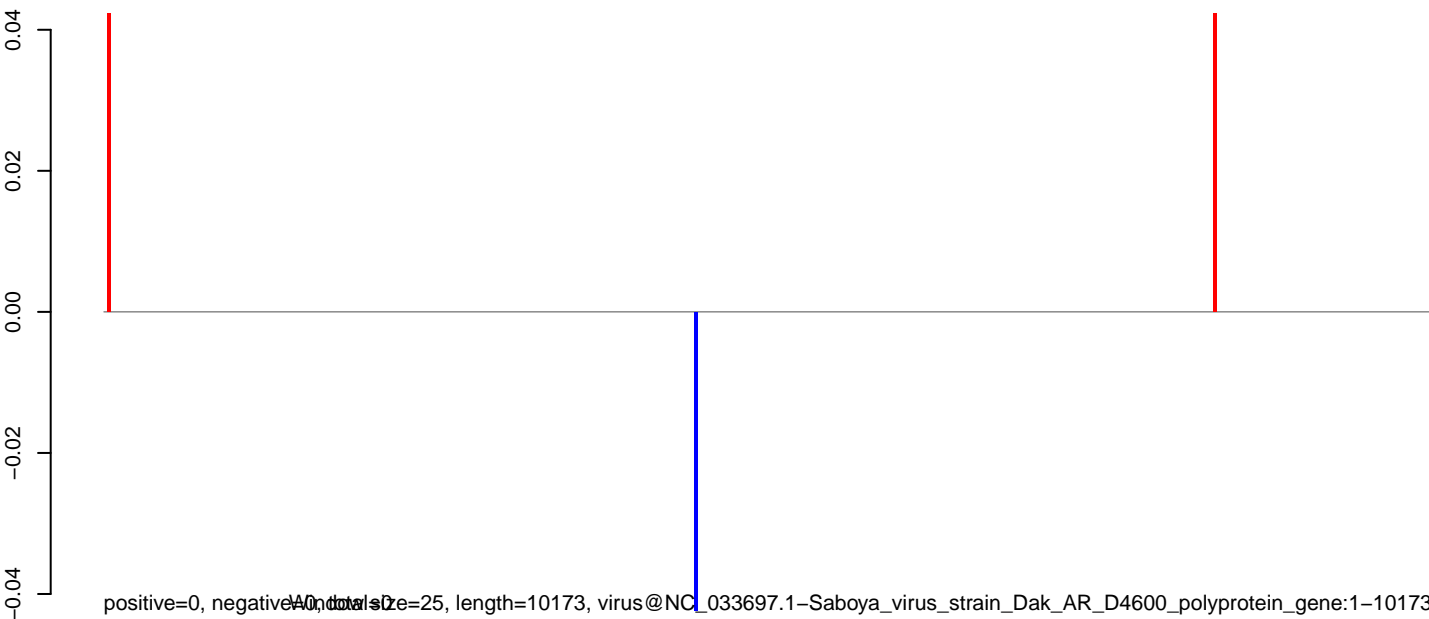
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

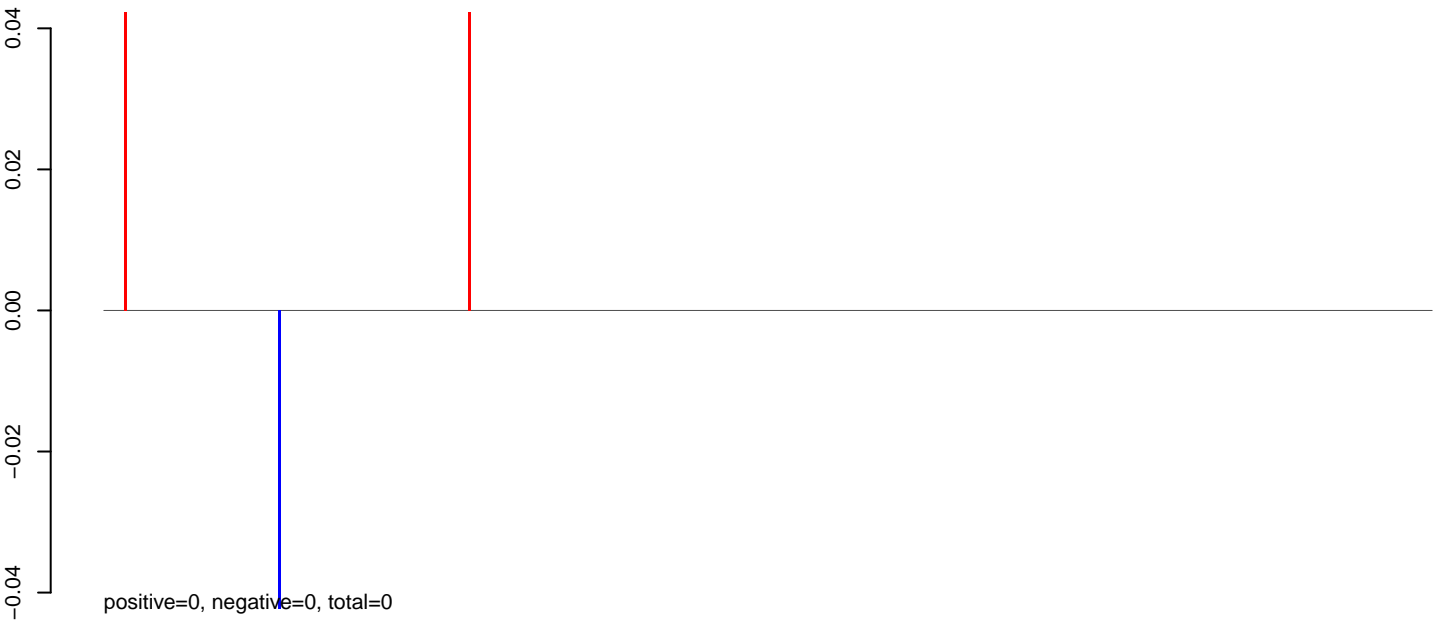


AeAlbo_C636_cells.rep

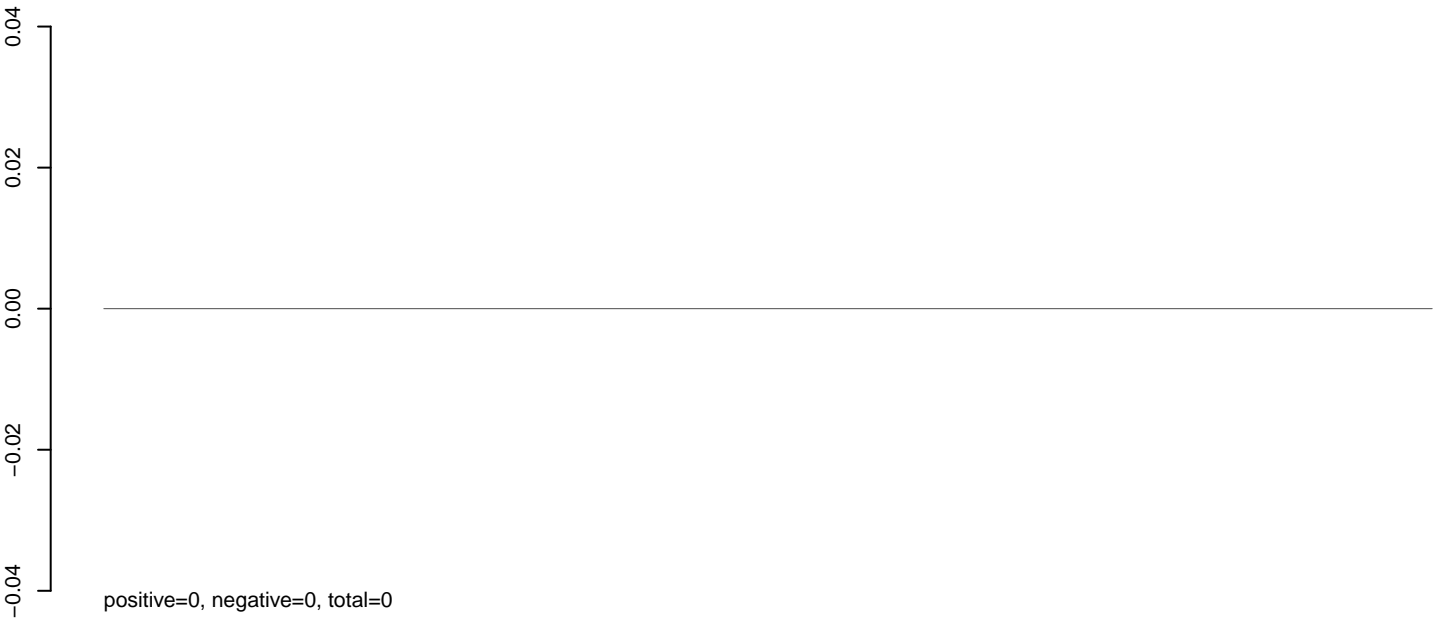


0 2000 4000 6000 8000 10000

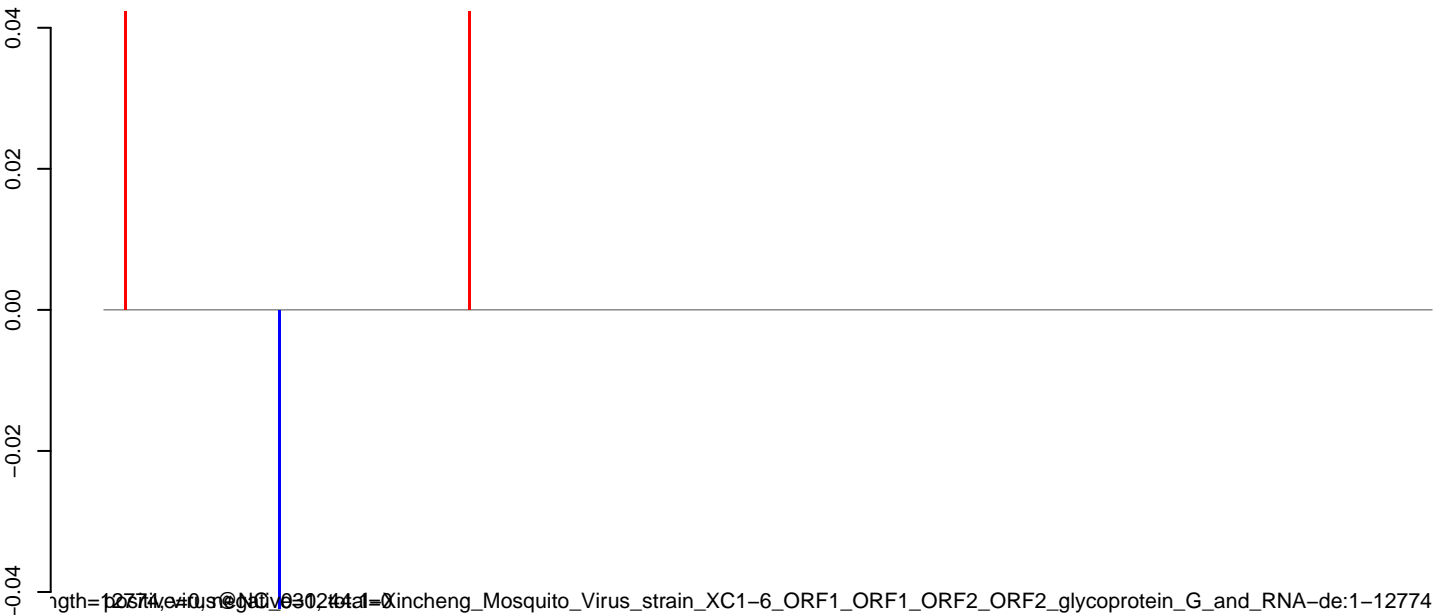
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

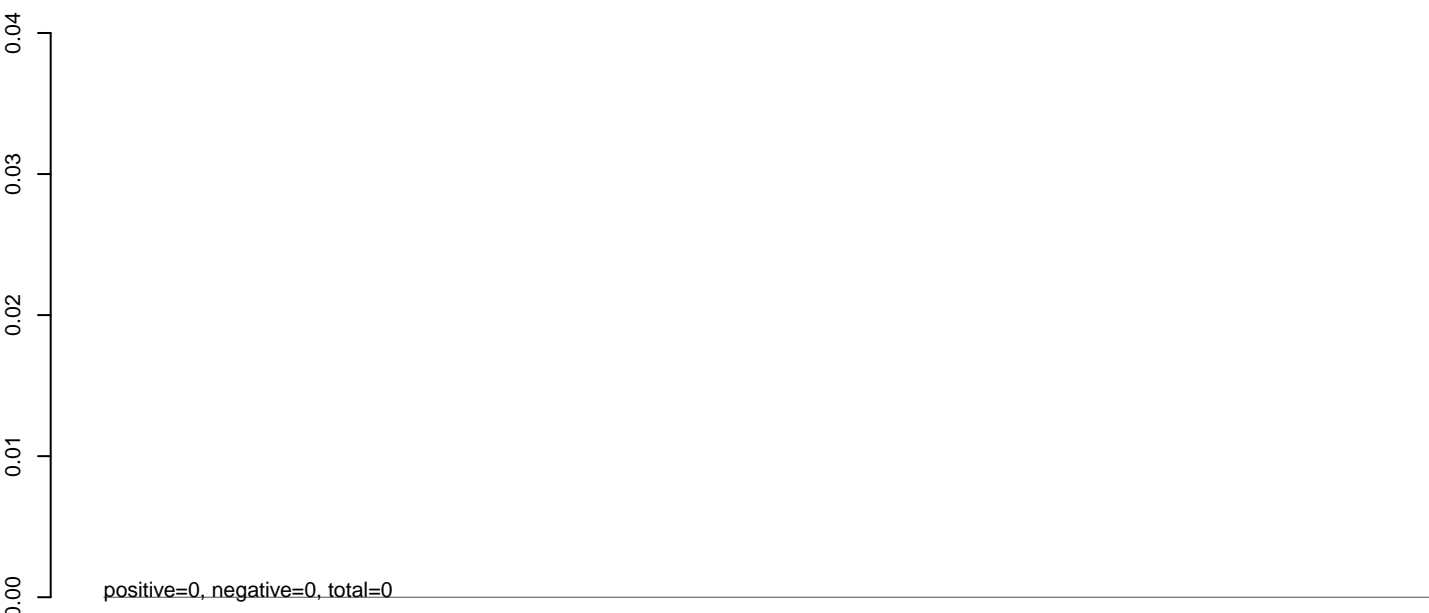


0 2000 4000 6000 8000 10000 12000

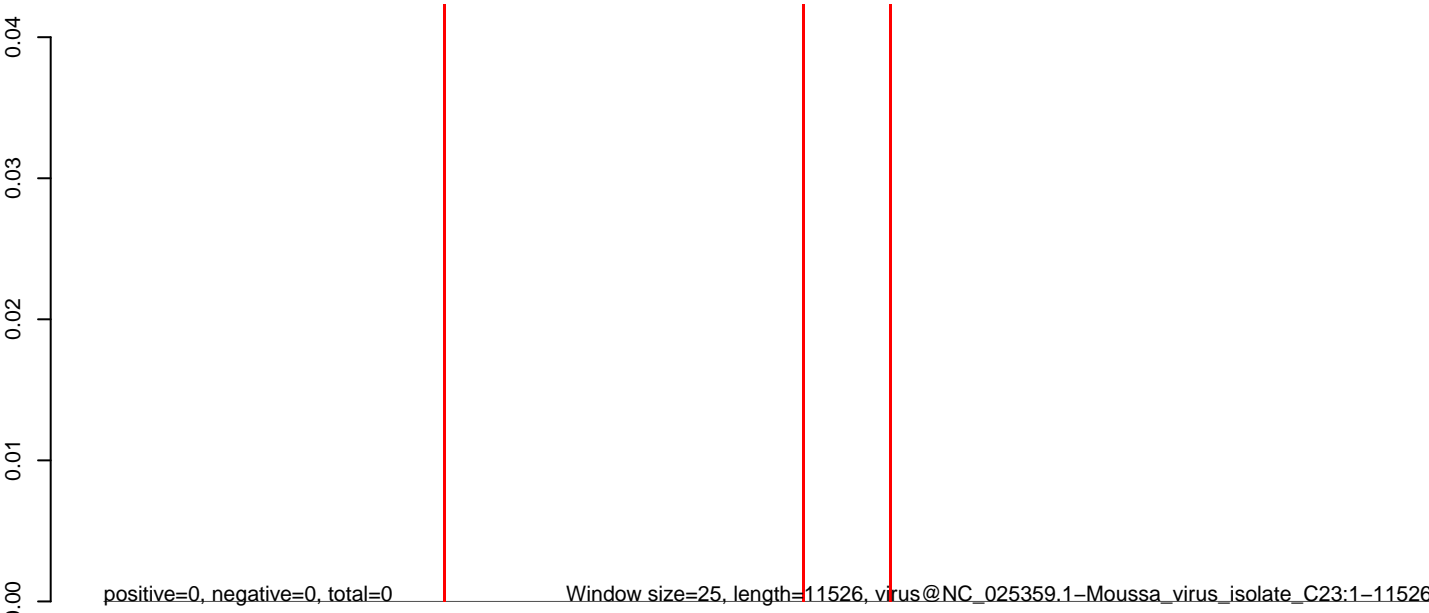
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

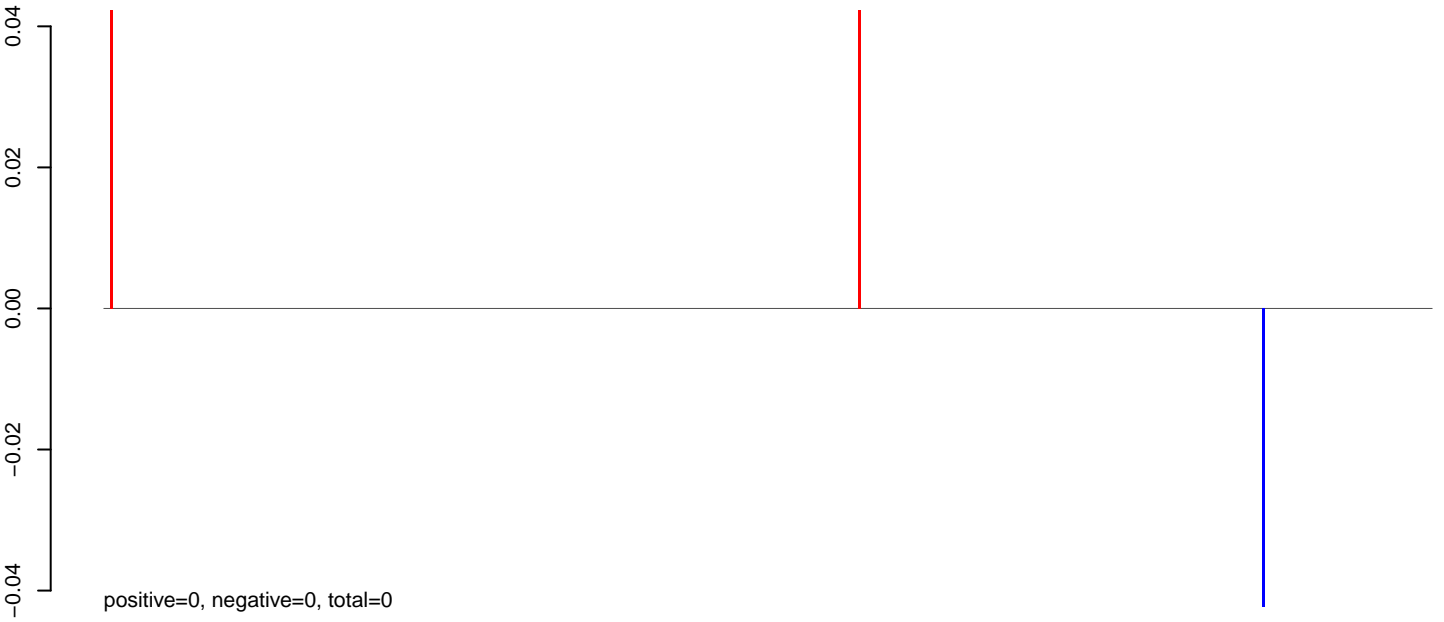


AeAlbo_C636_cells.rep

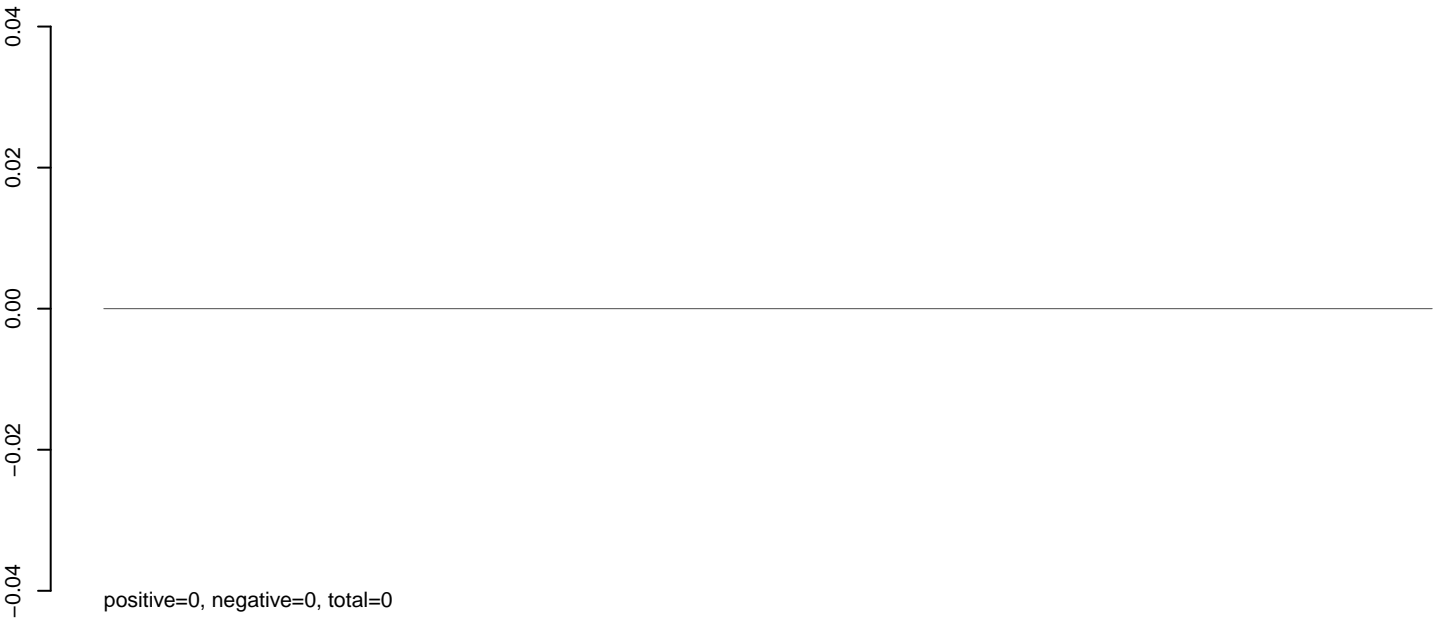


0 2000 4000 6000 8000 10000 12000

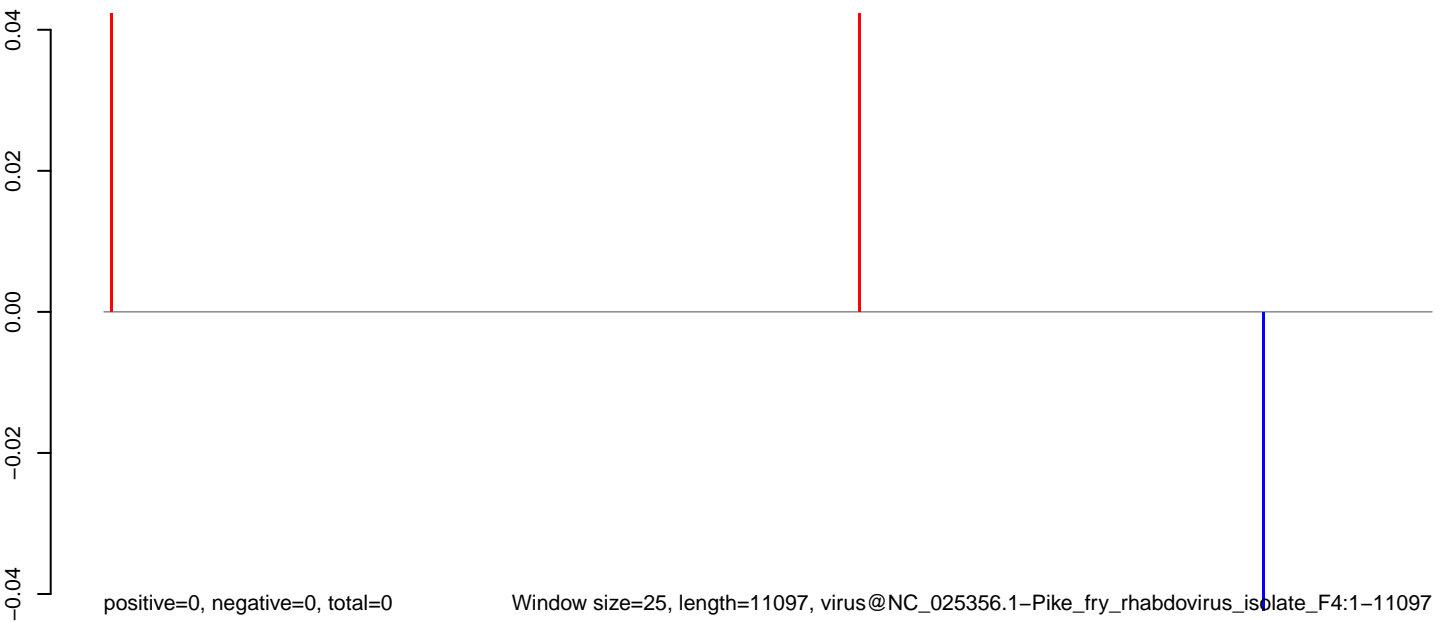
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



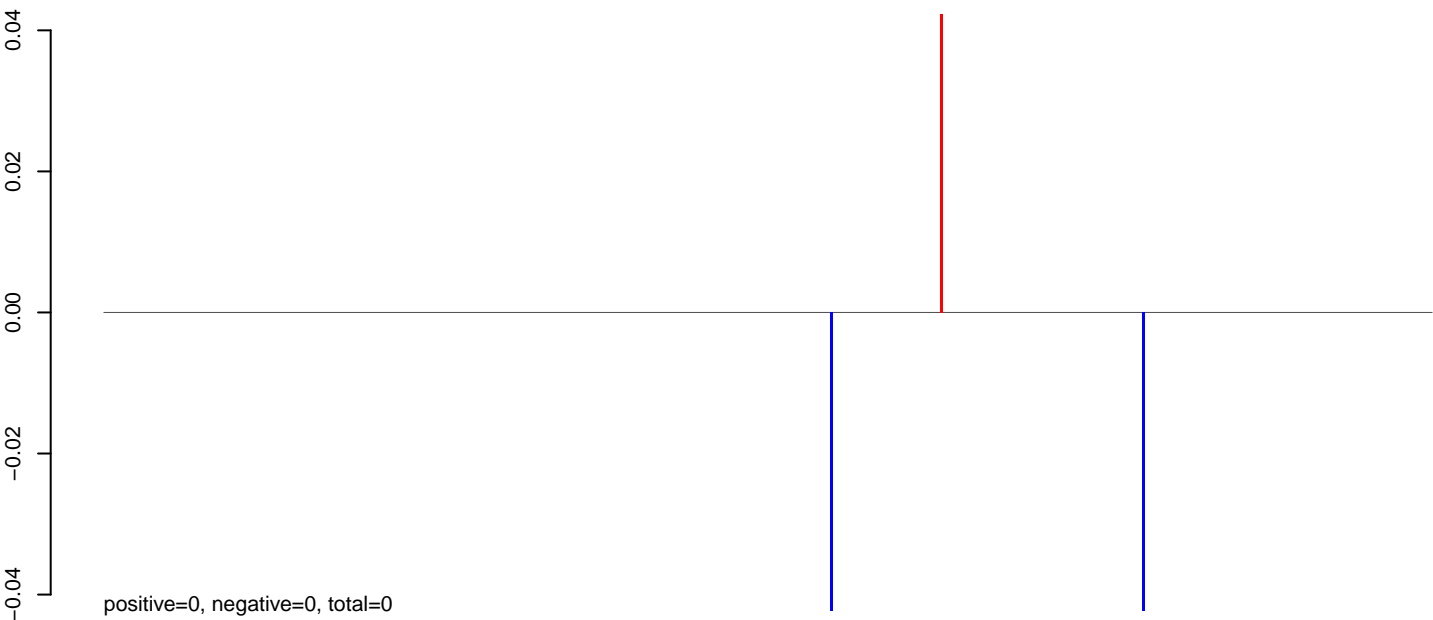
AeAlbo_C636_cells.rep



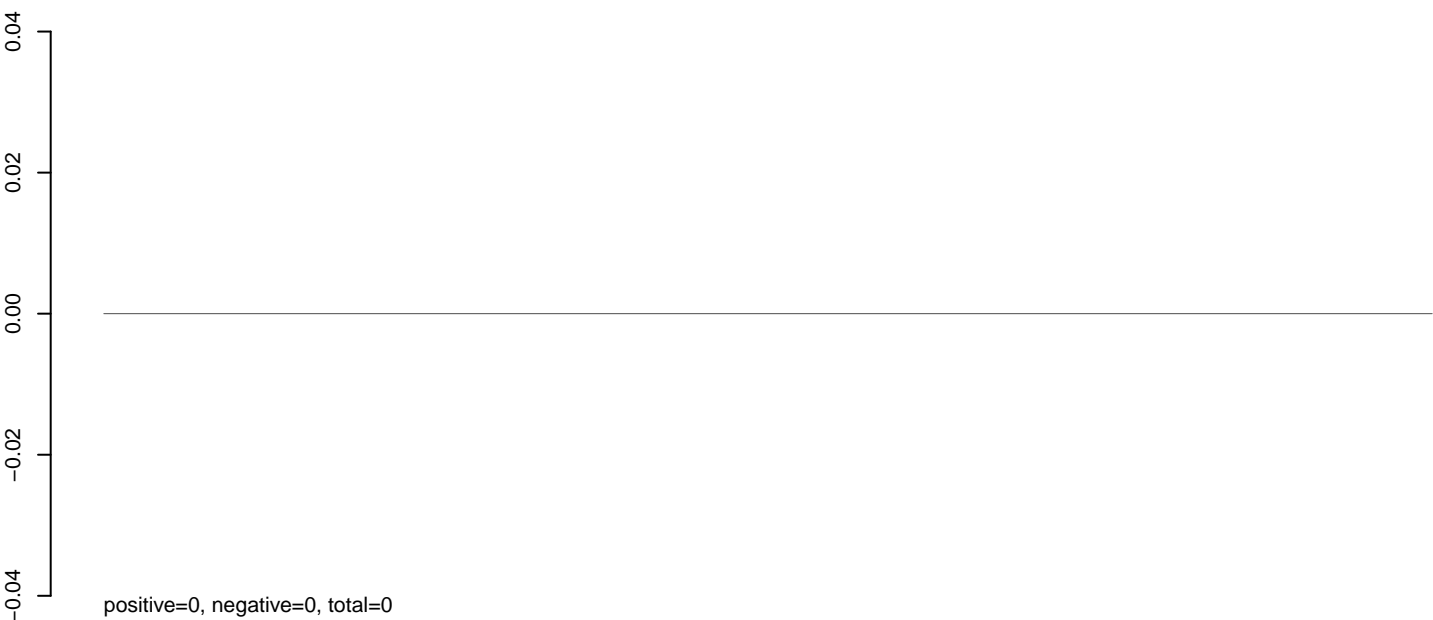
Window size=25, length=11097, virus@NC_025356.1-Pike_fry_rhabdovirus_isolate_F4:1-11097

0 2000 4000 6000 8000 10000

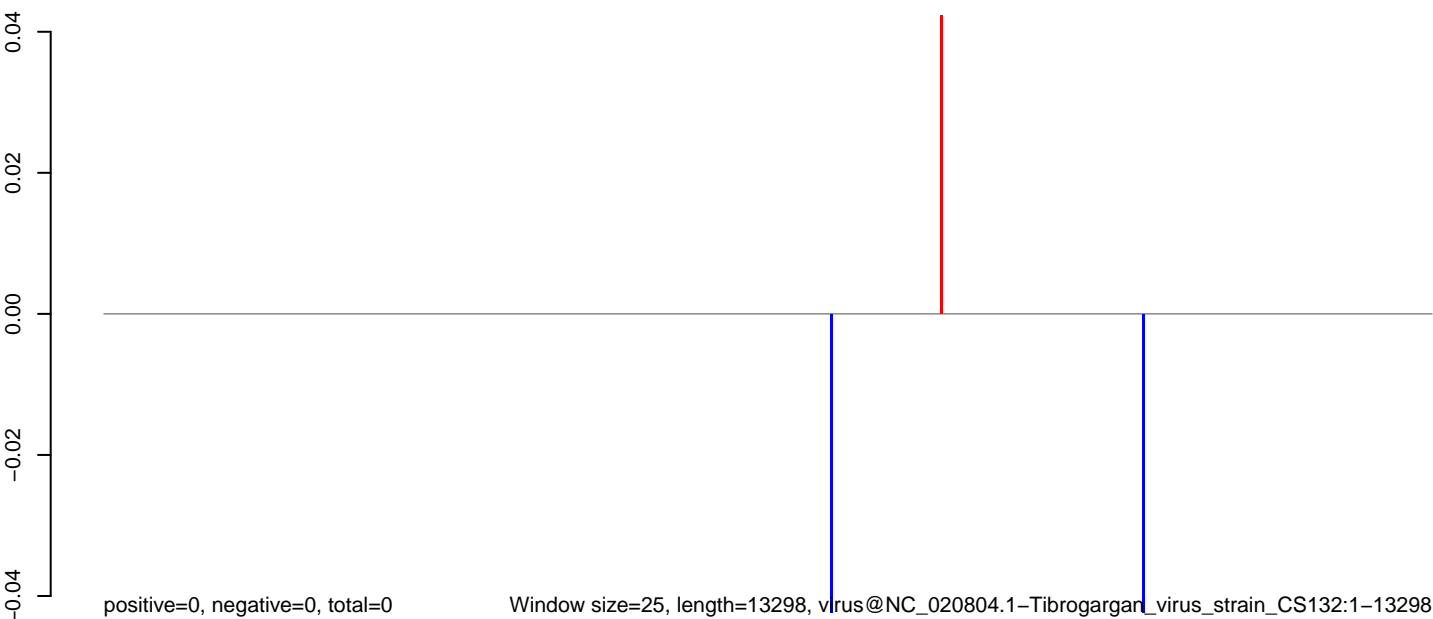
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



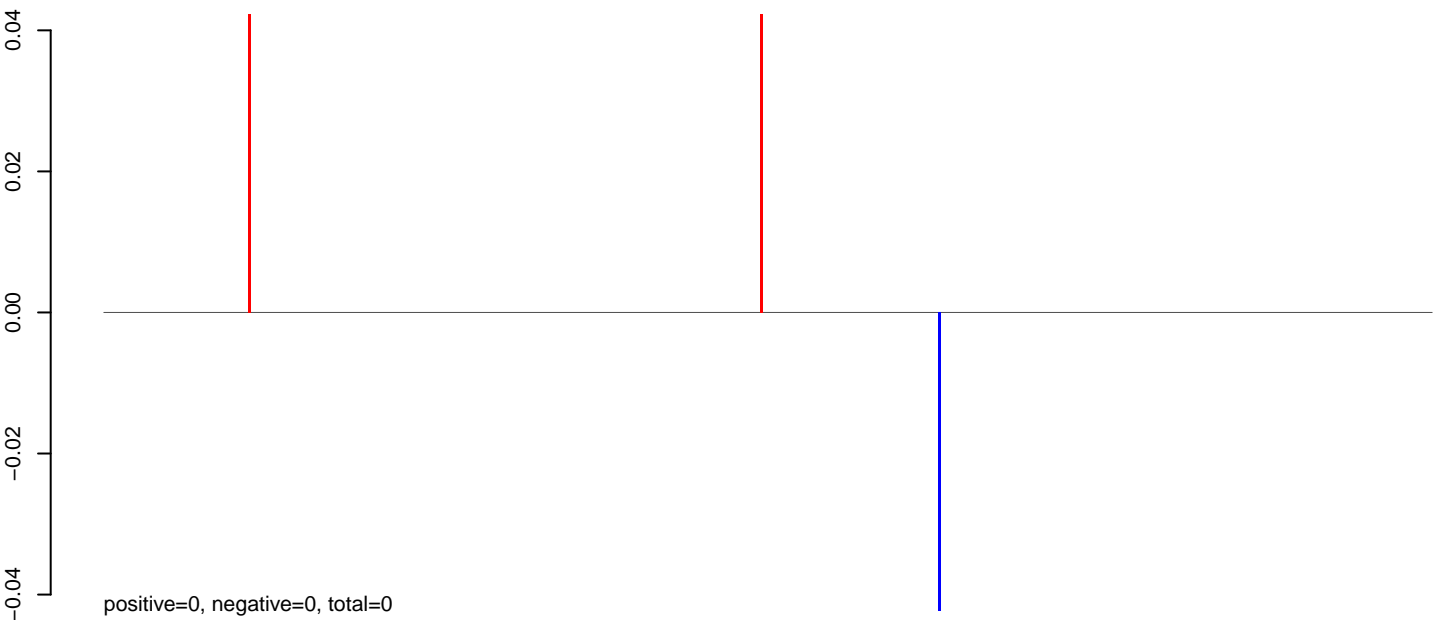
AeAlbo_C636_cells.rep



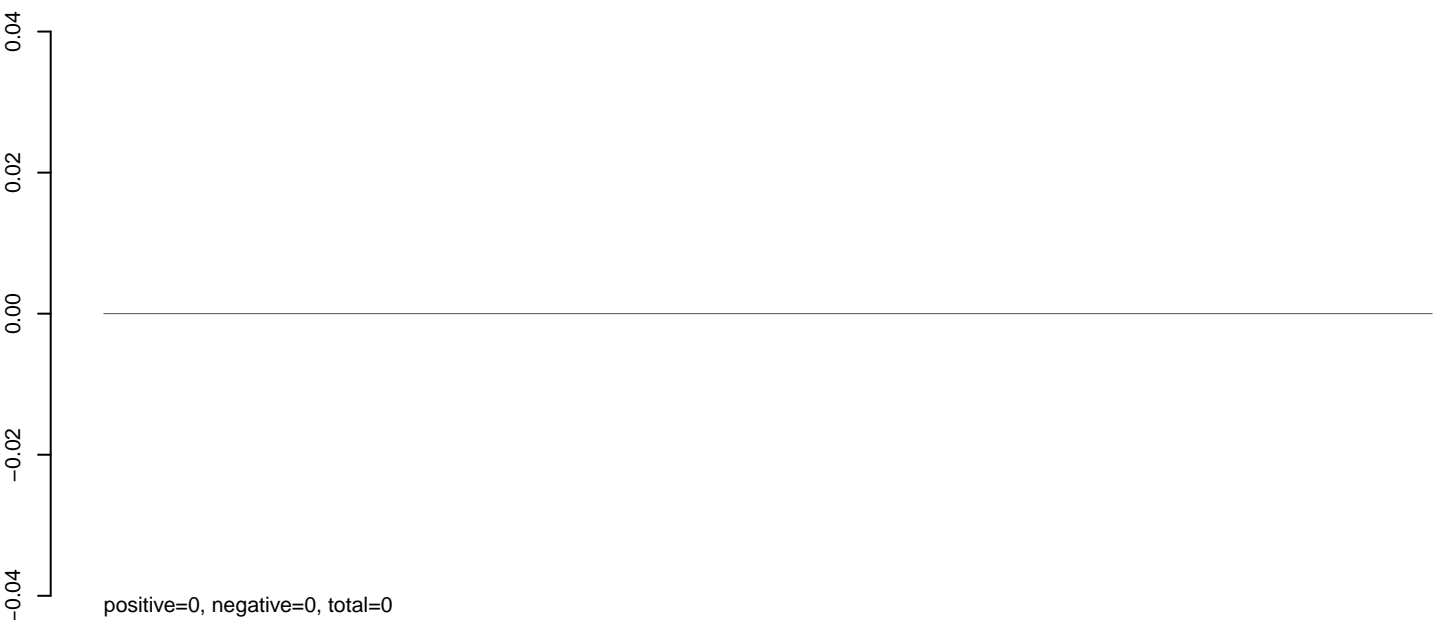
Window size=25, length=13298, virus@NC_020804.1-Tibrogargan_virus_strain_CS132:1-13298

0 2000 4000 6000 8000 10000 12000

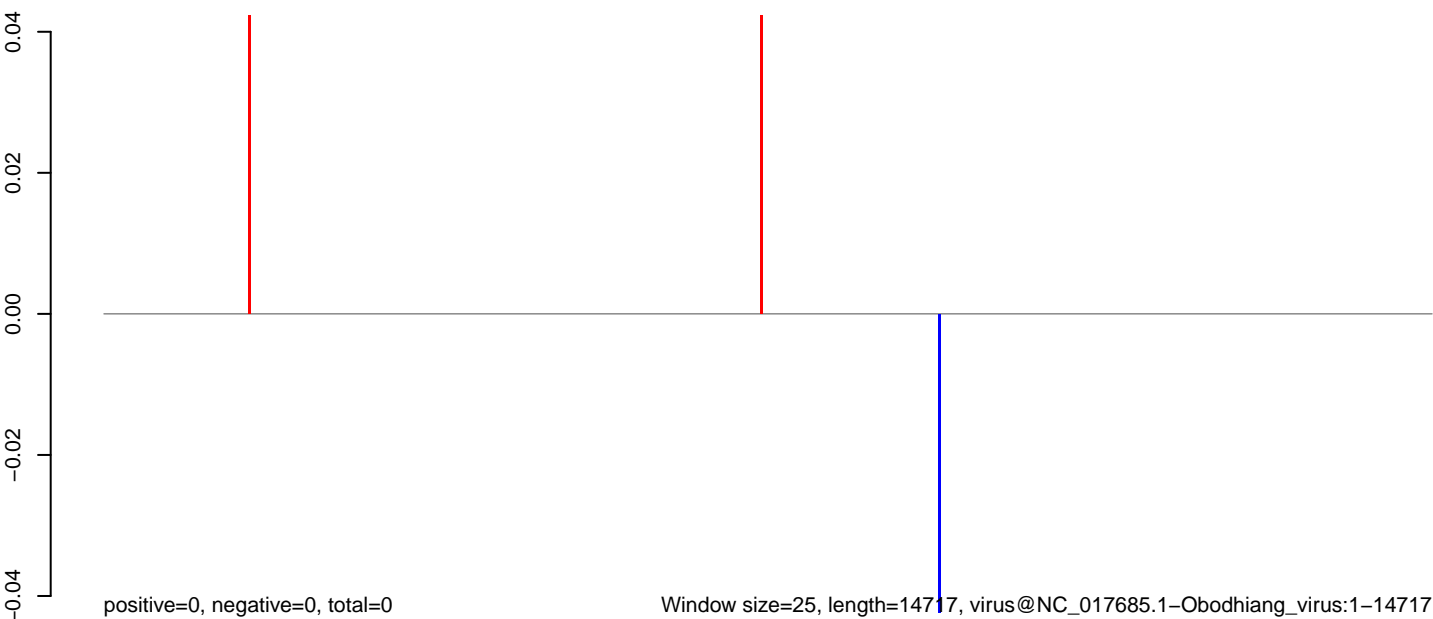
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



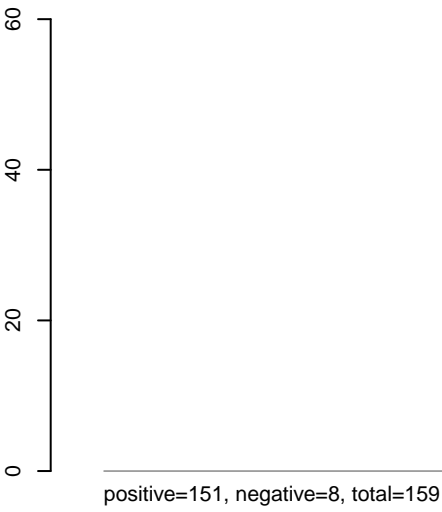
AeAlbo_C636_cells.rep



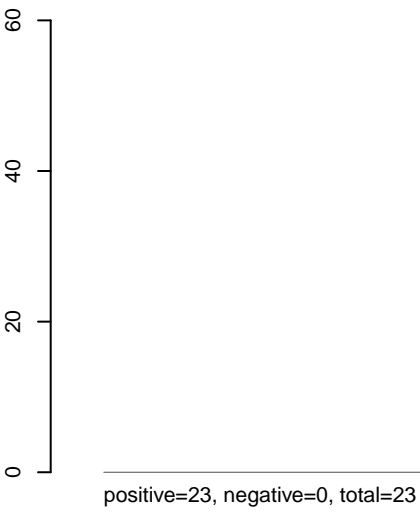
Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

0 5000 10000 15000

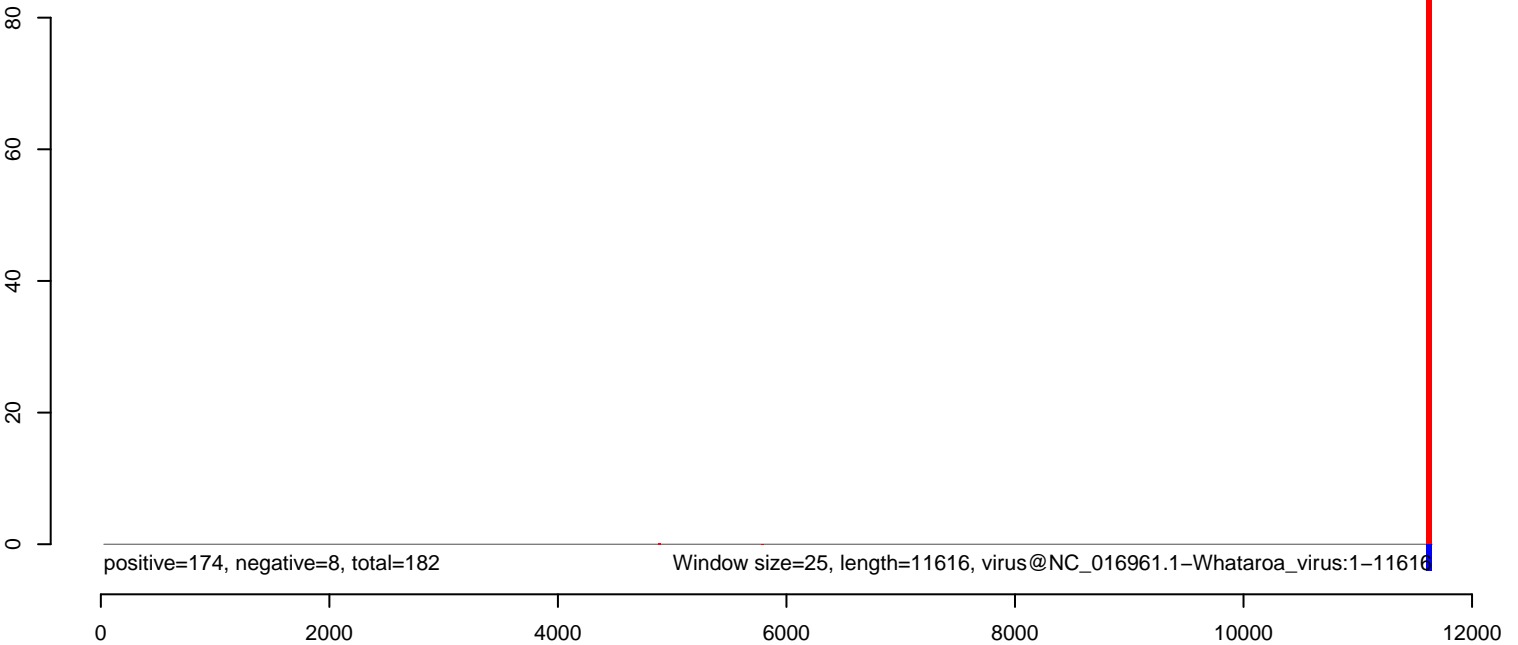
AeAlbo_C636_cells.18_23.rep



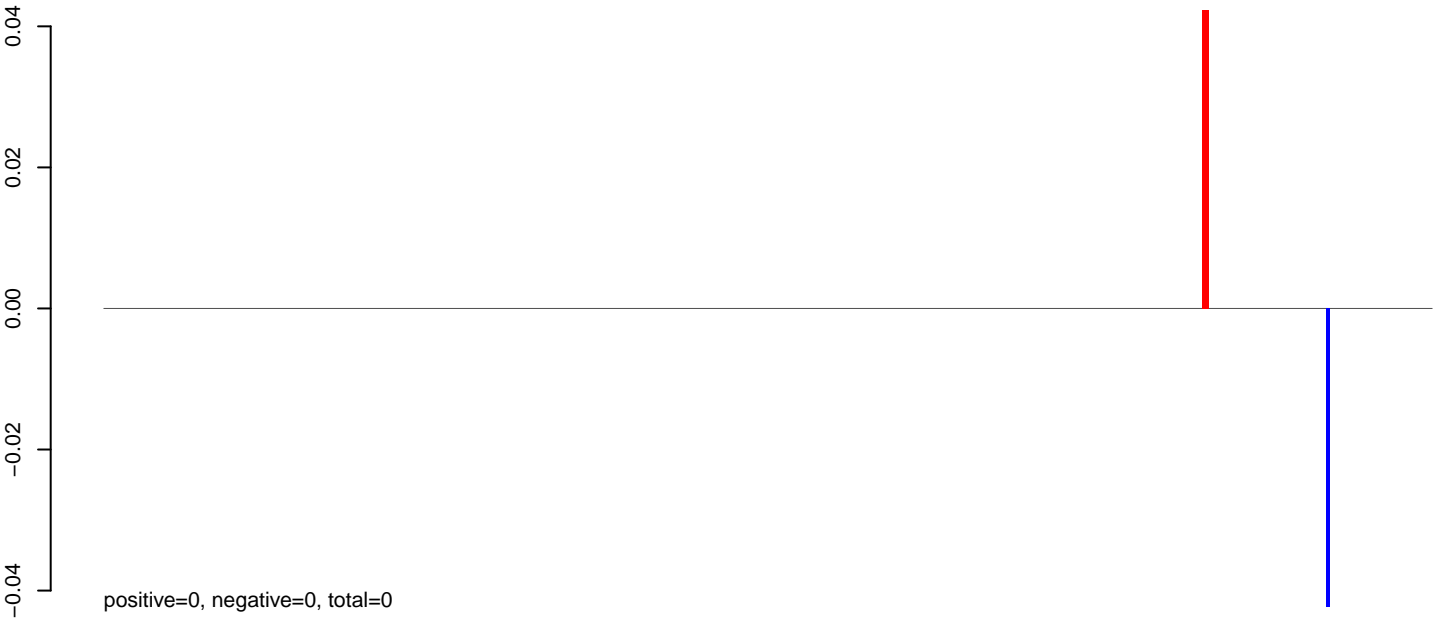
AeAlbo_C636_cells.24_35.rep



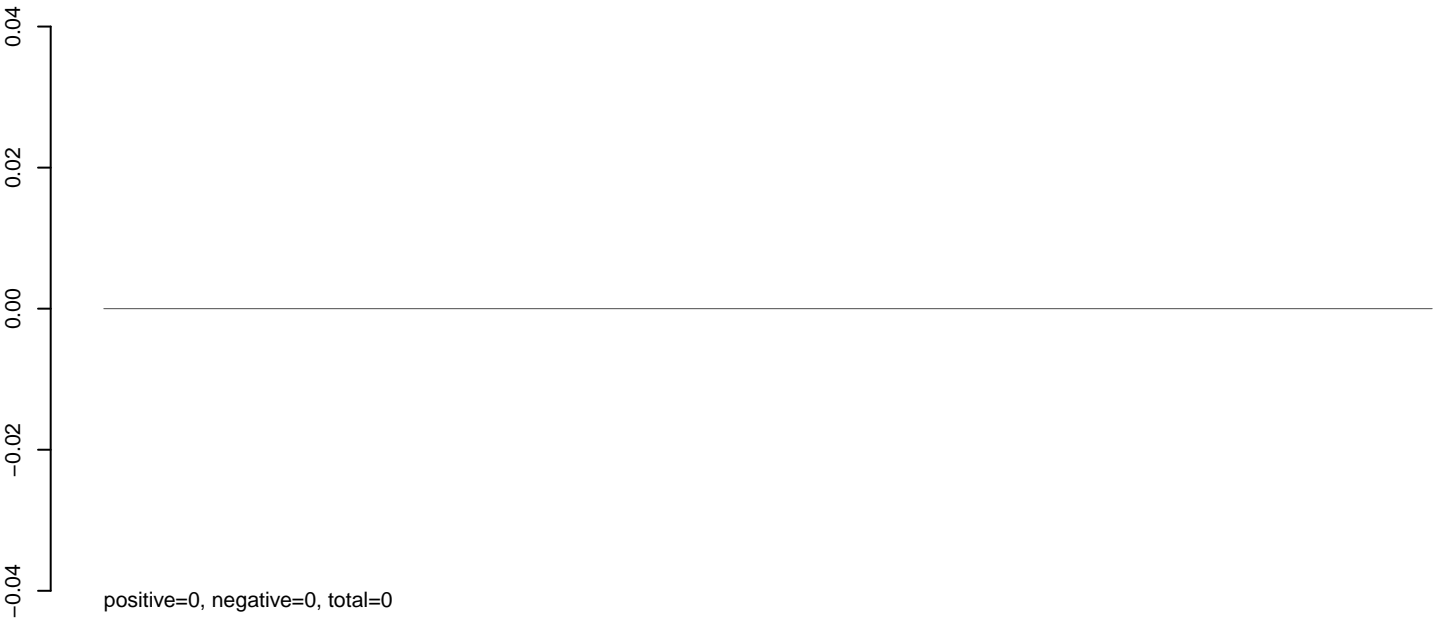
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



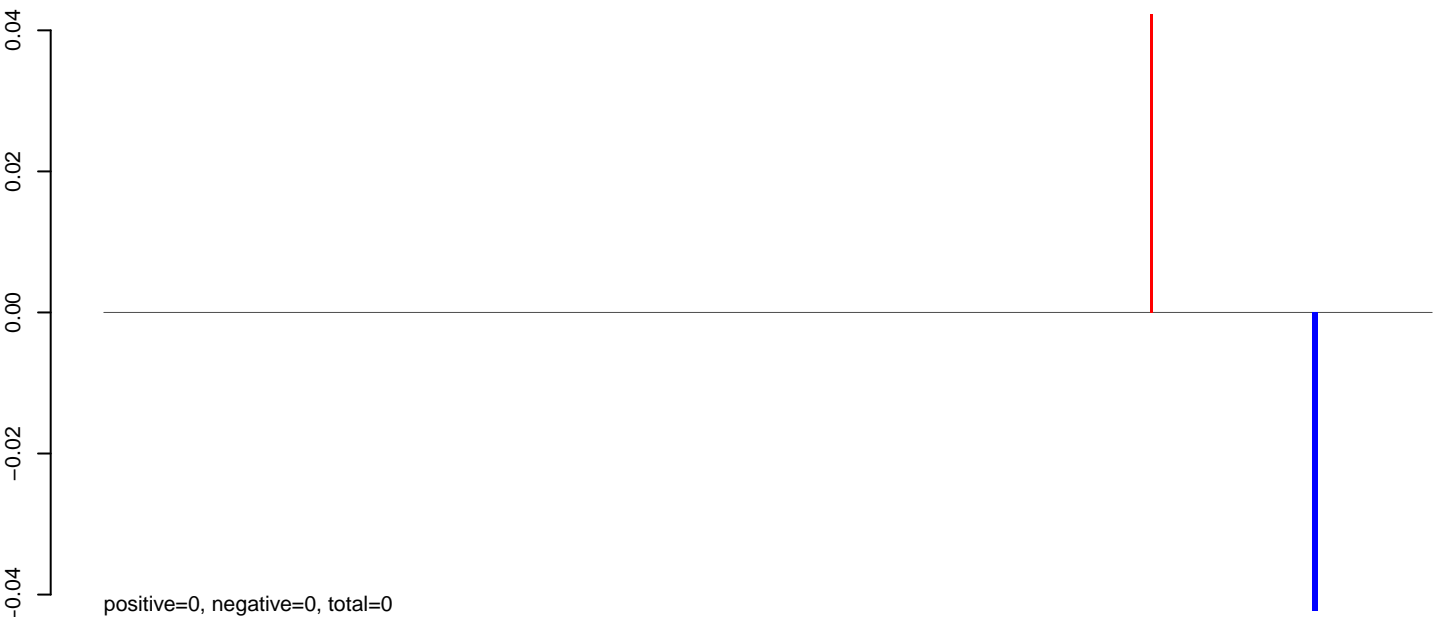
AeAlbo_C636_cells.rep



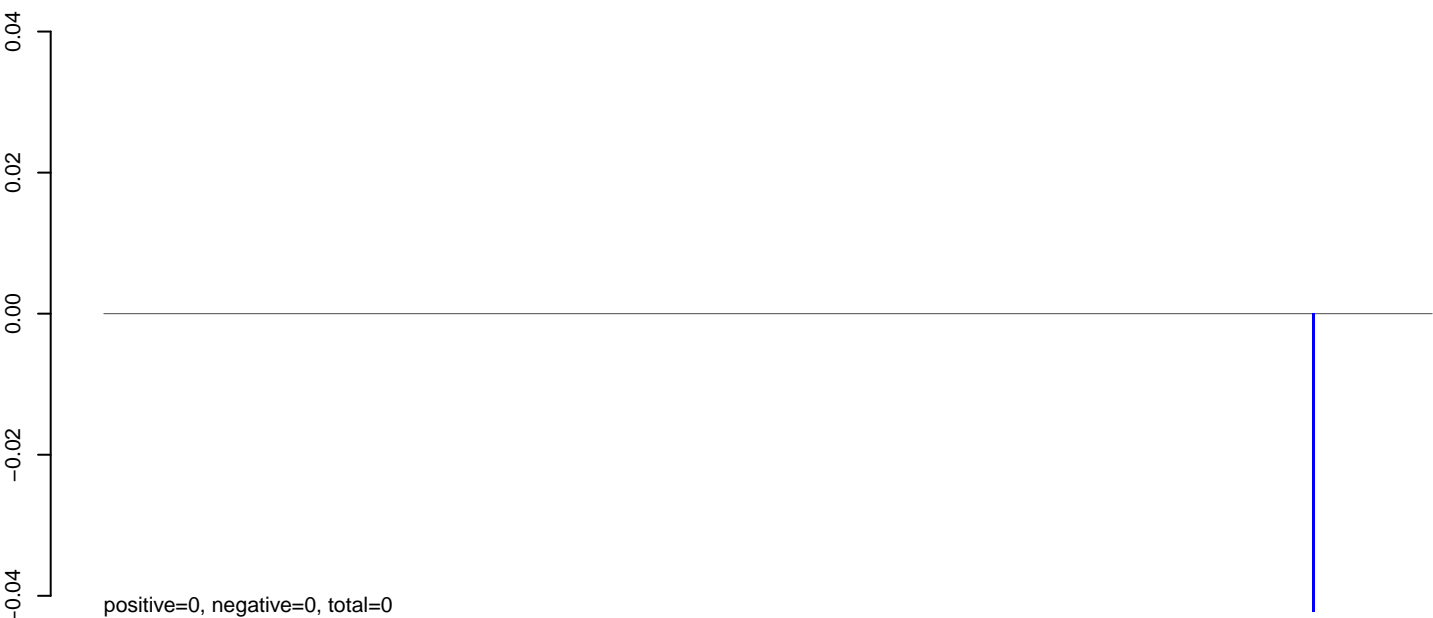
Windows size=25, length=10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

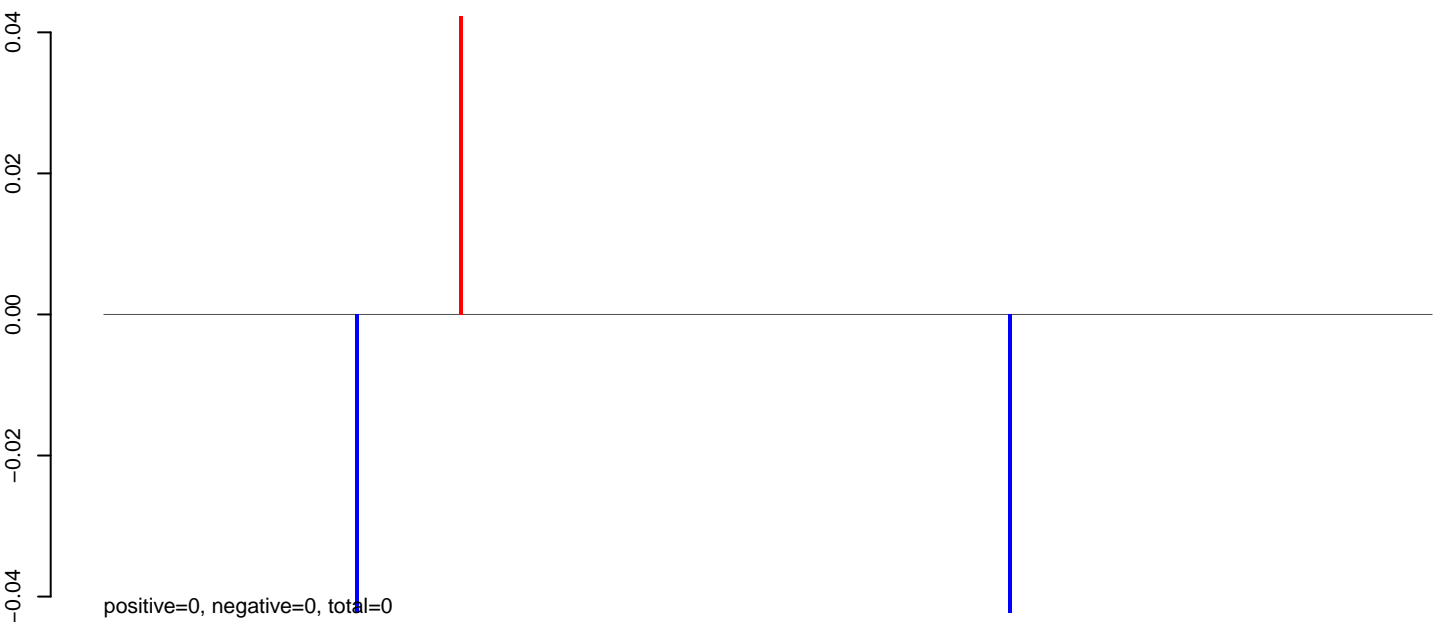


AeAlbo_C636_cells.rep

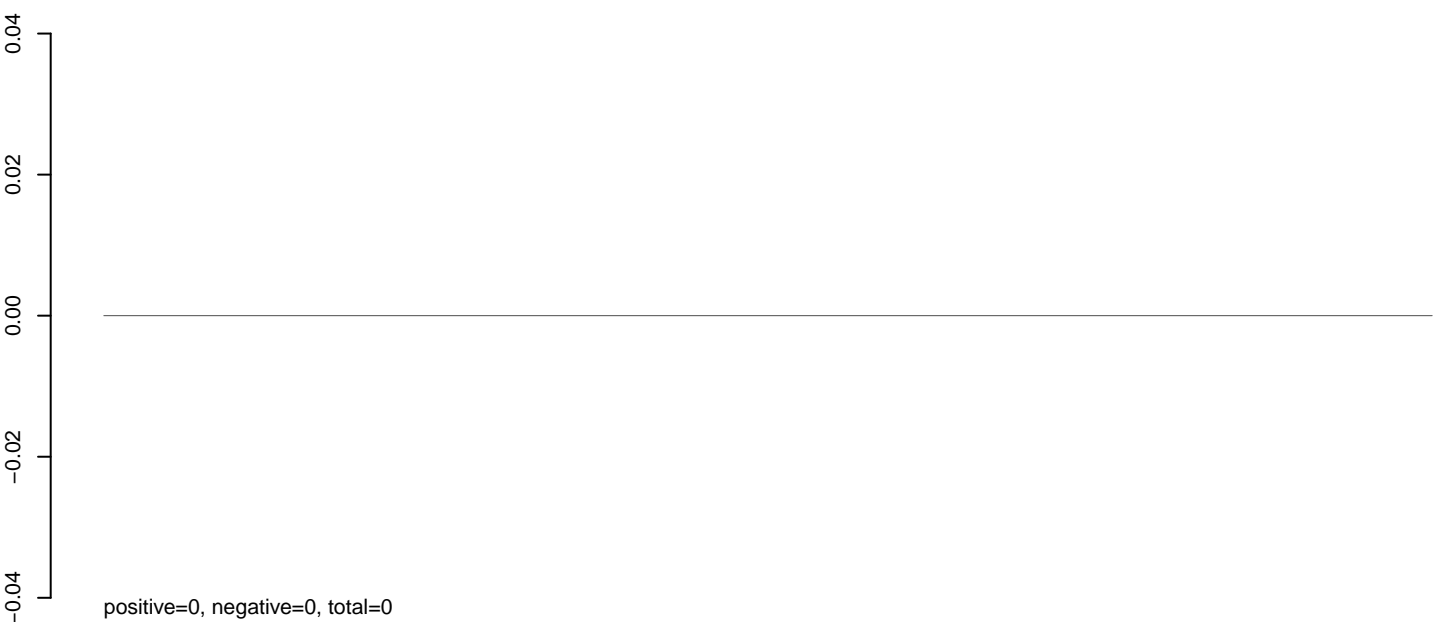


0 2000 4000 6000 8000 10000

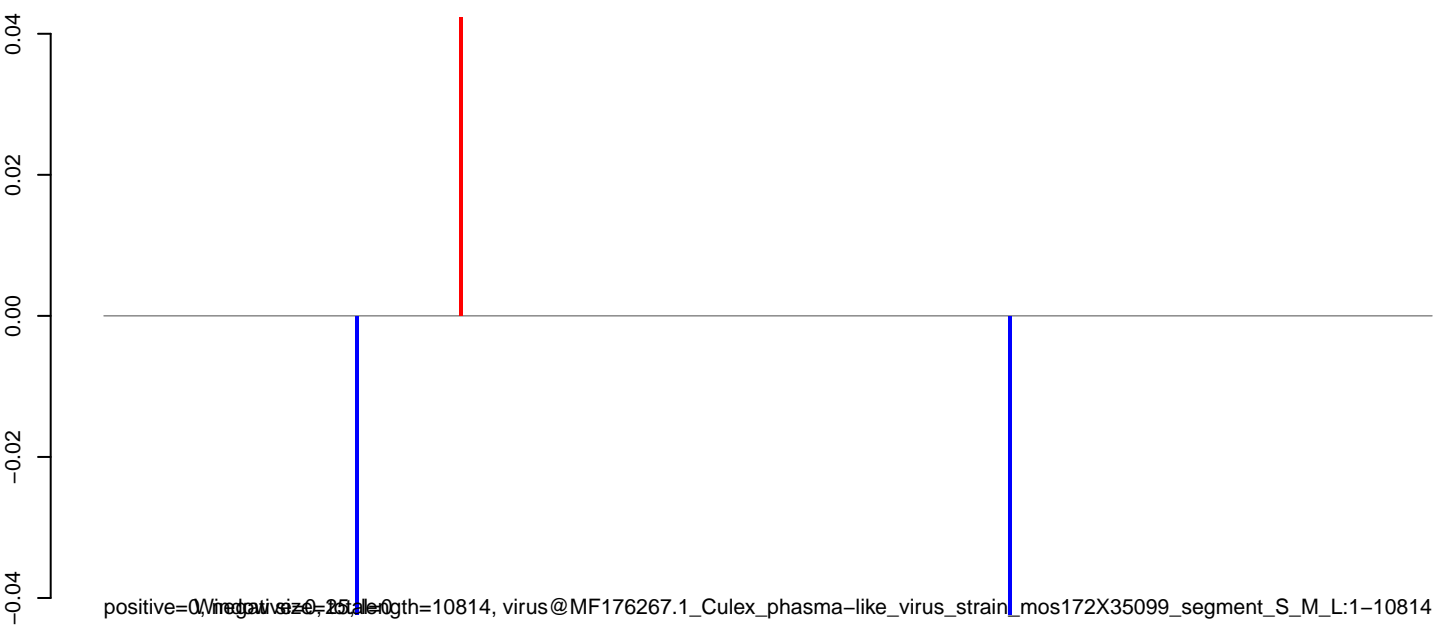
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

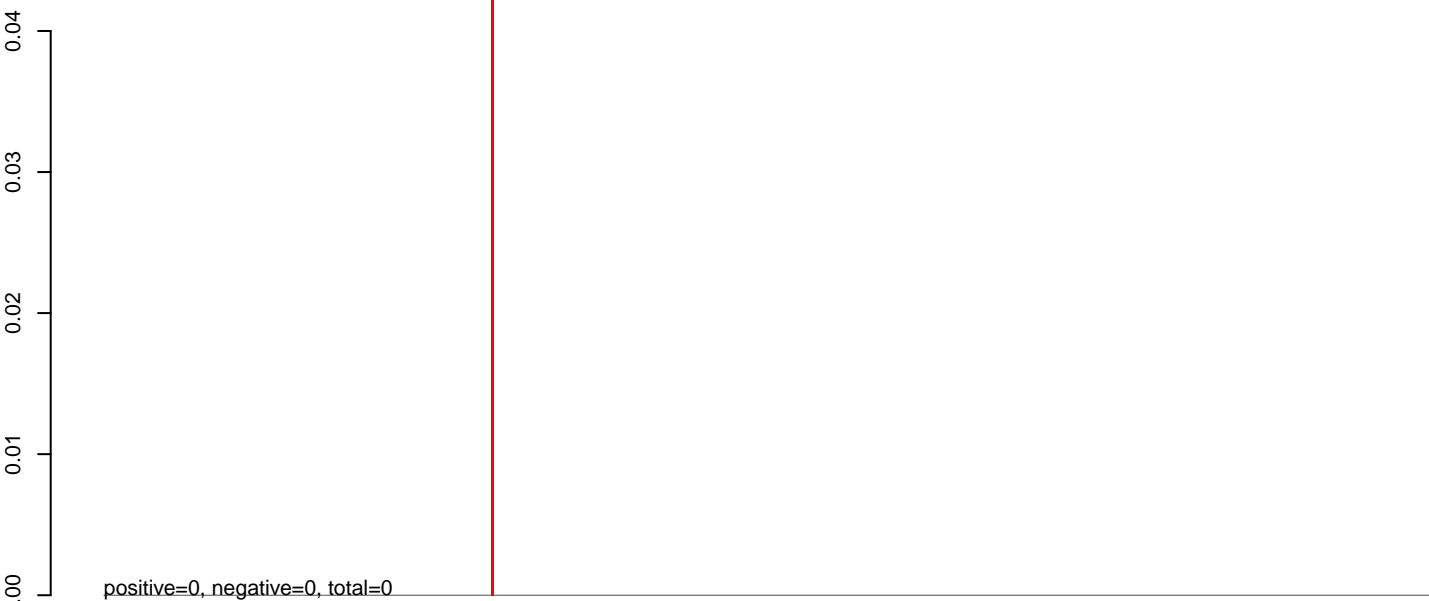


0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



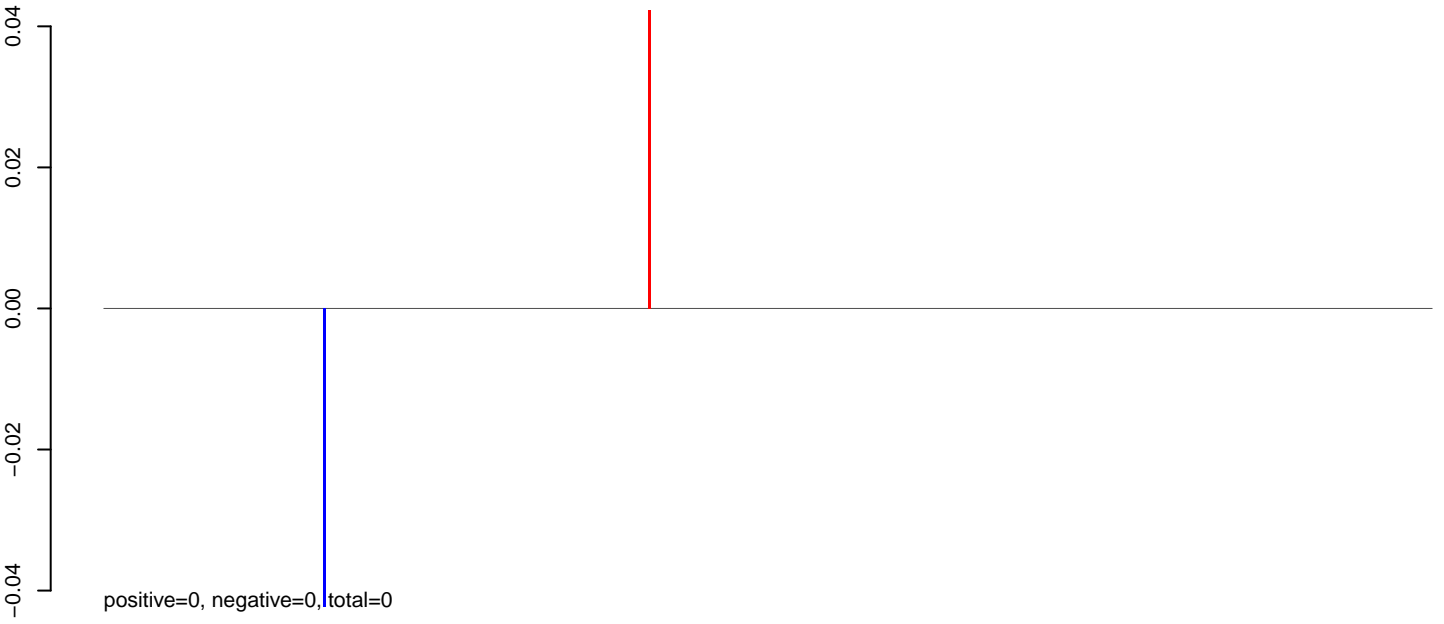
AeAlbo_C636_cells.24_35.rep



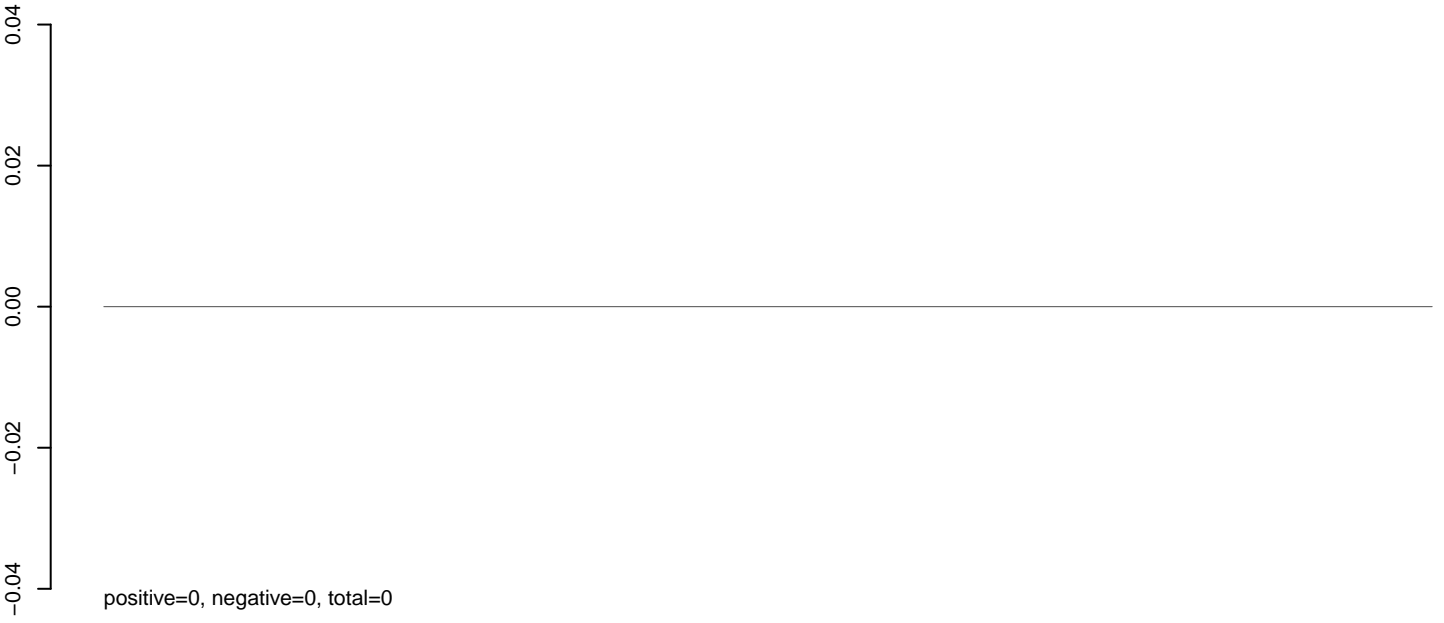
AeAlbo_C636_cells.rep



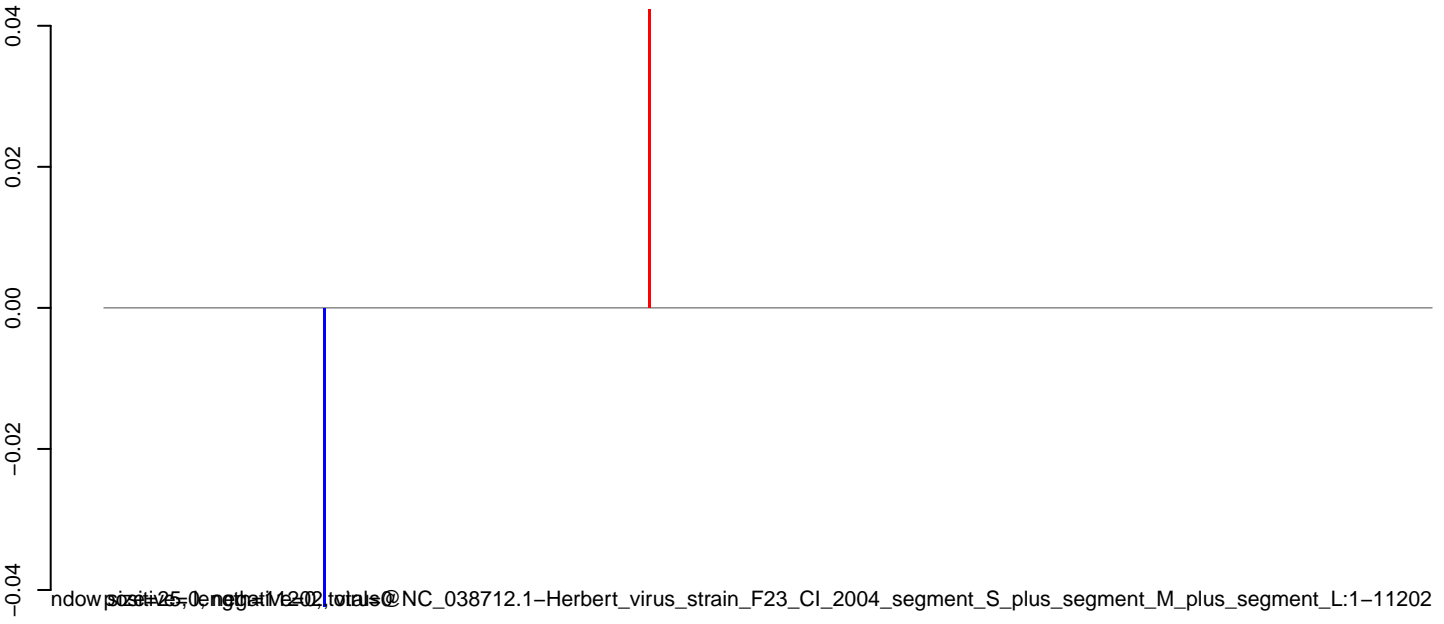
AeAlbo_C636_cells.18_23.rep



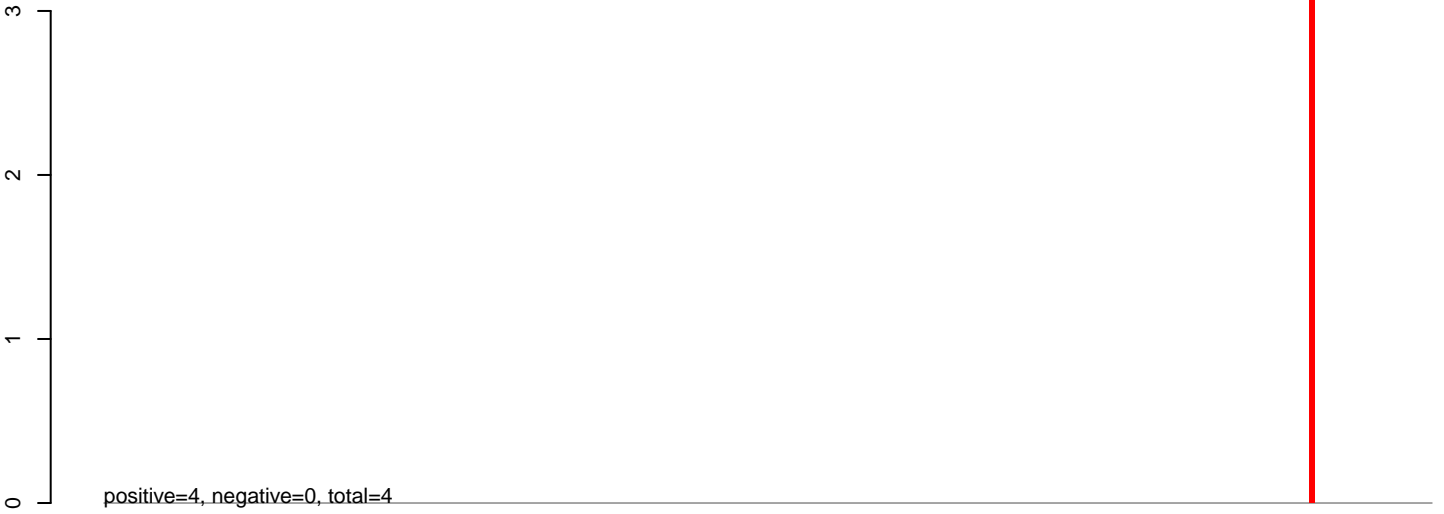
AeAlbo_C636_cells.24_35.rep



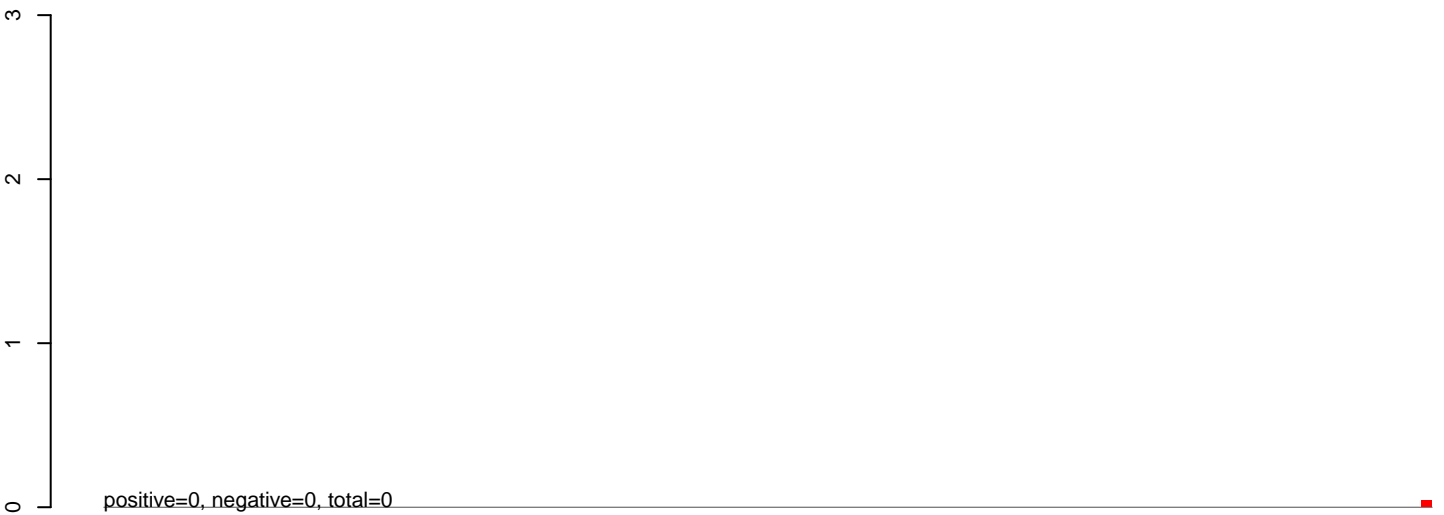
AeAlbo_C636_cells.rep



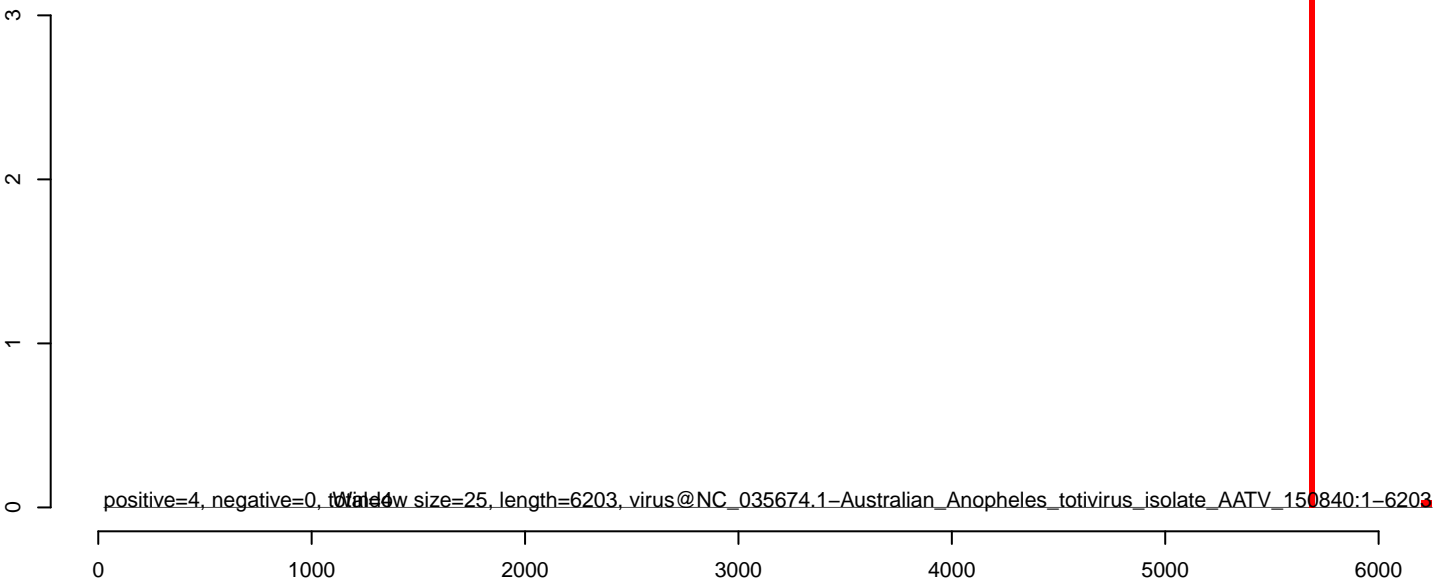
AeAlbo_C636_cells.18_23.rep



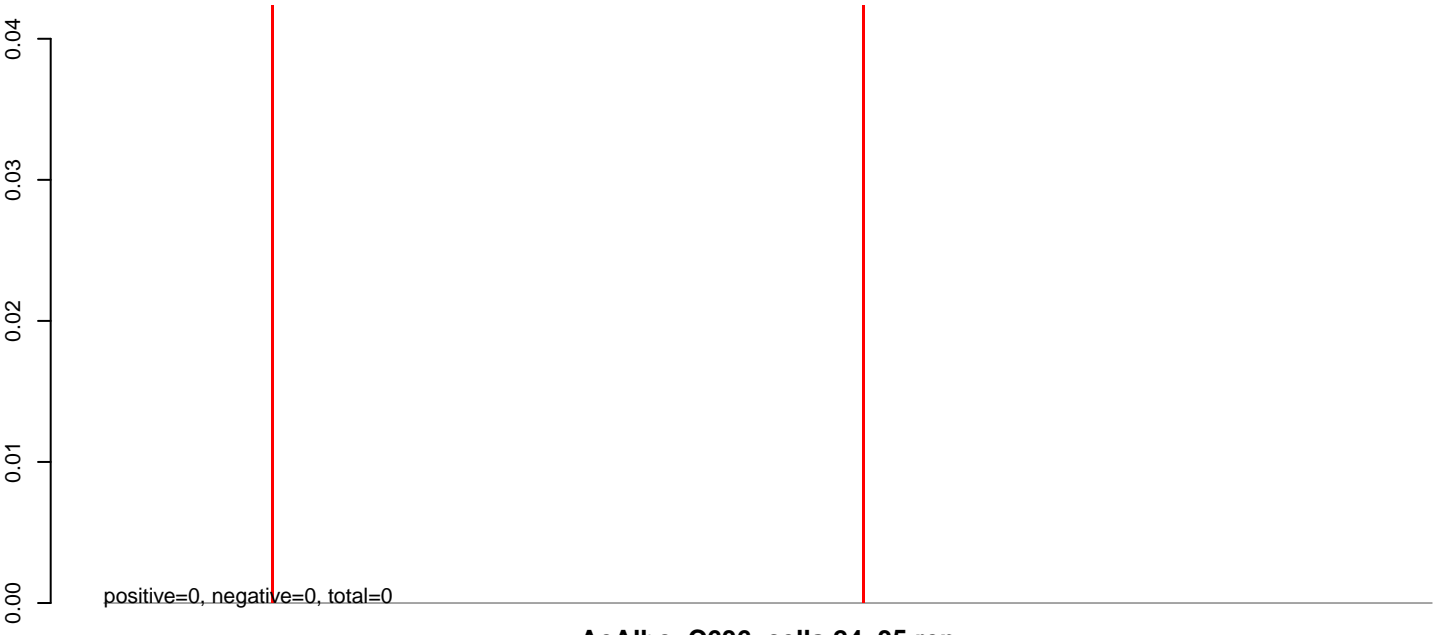
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



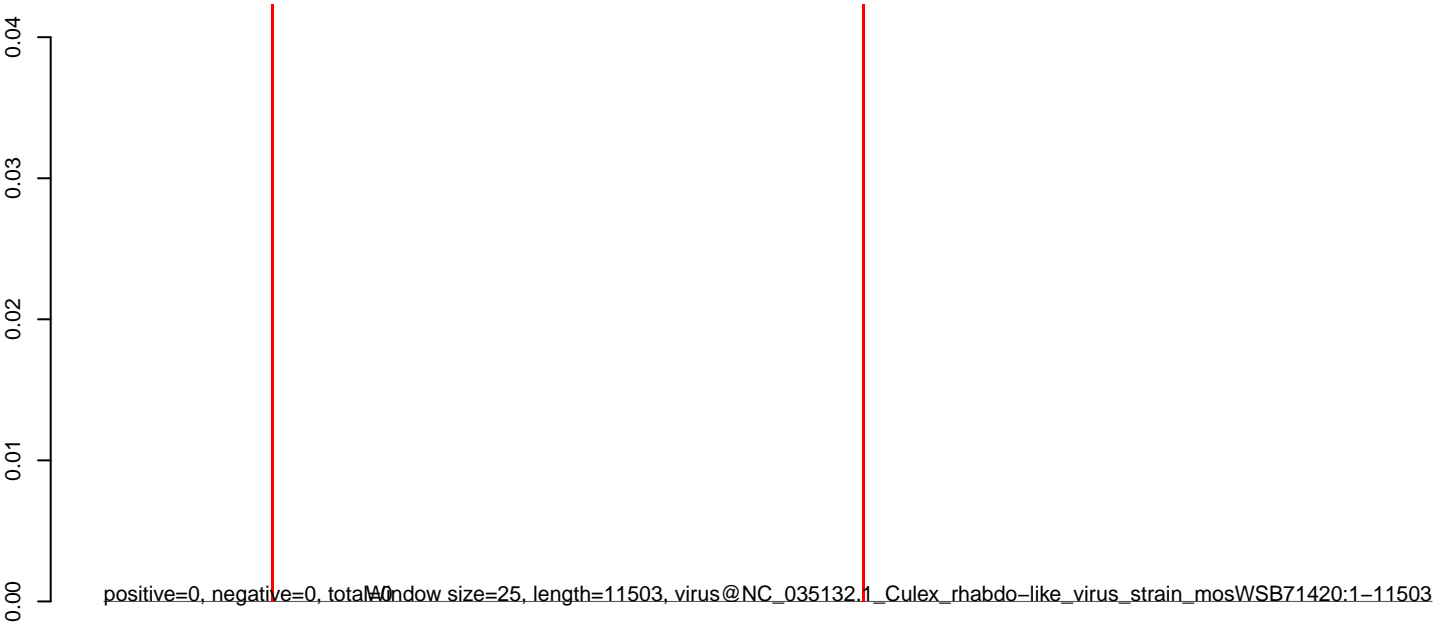
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

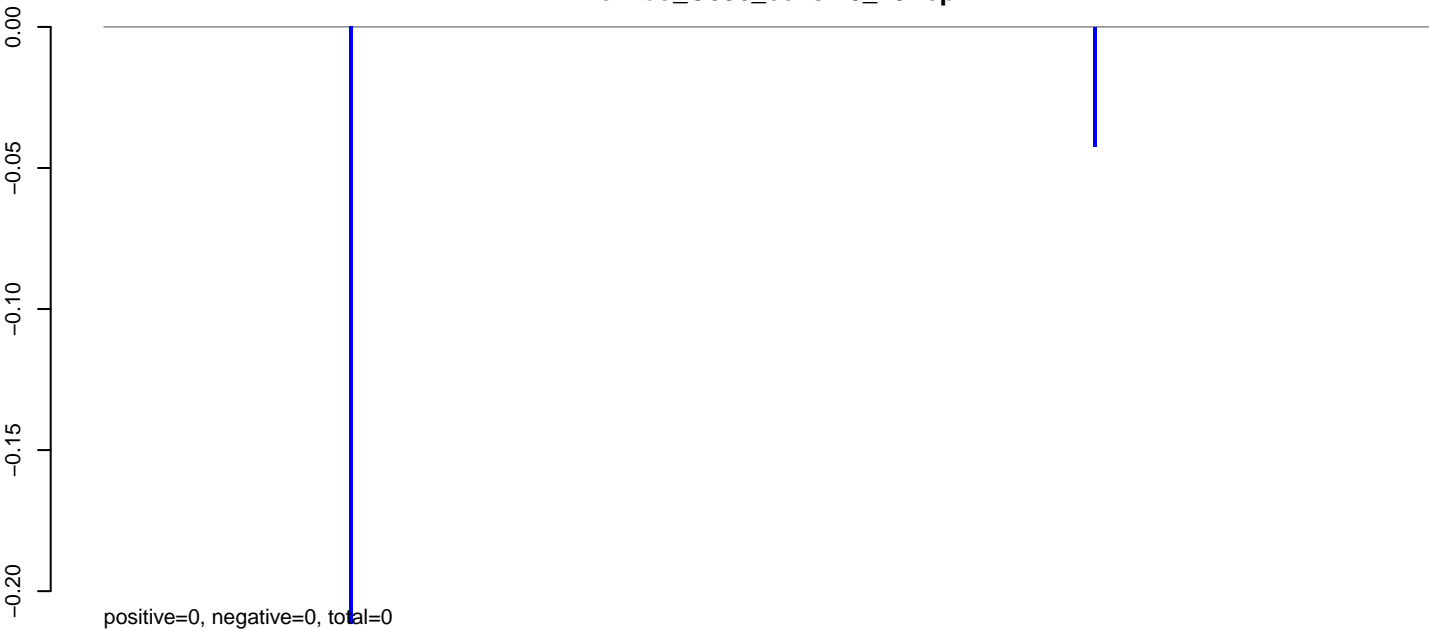


AeAlbo_C636_cells.rep

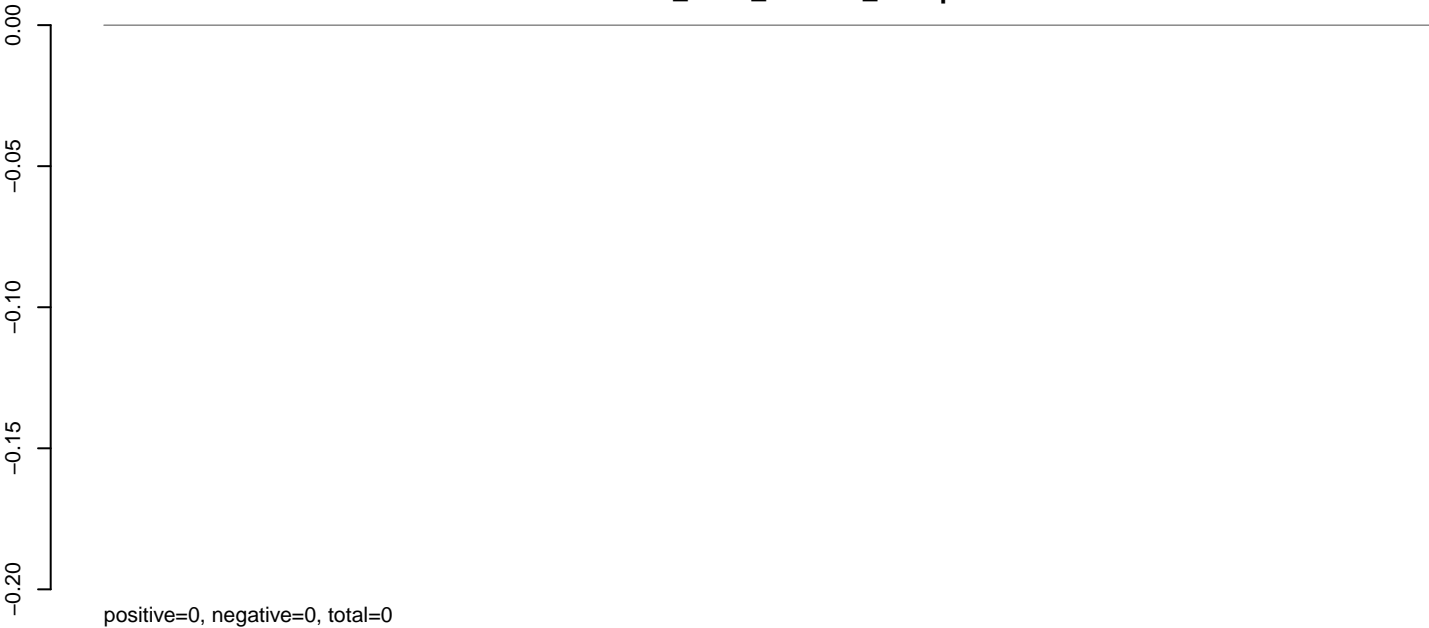


Window size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503

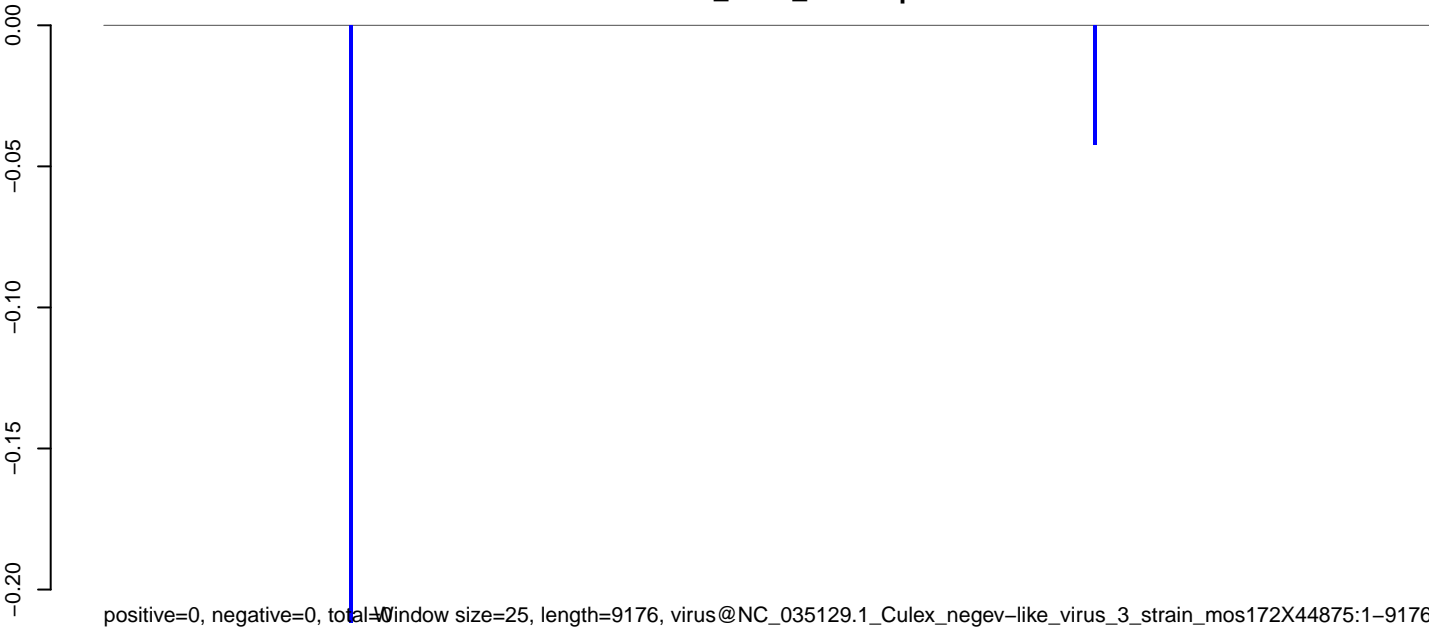
AeAlbo_C636_cells.18_23.rep



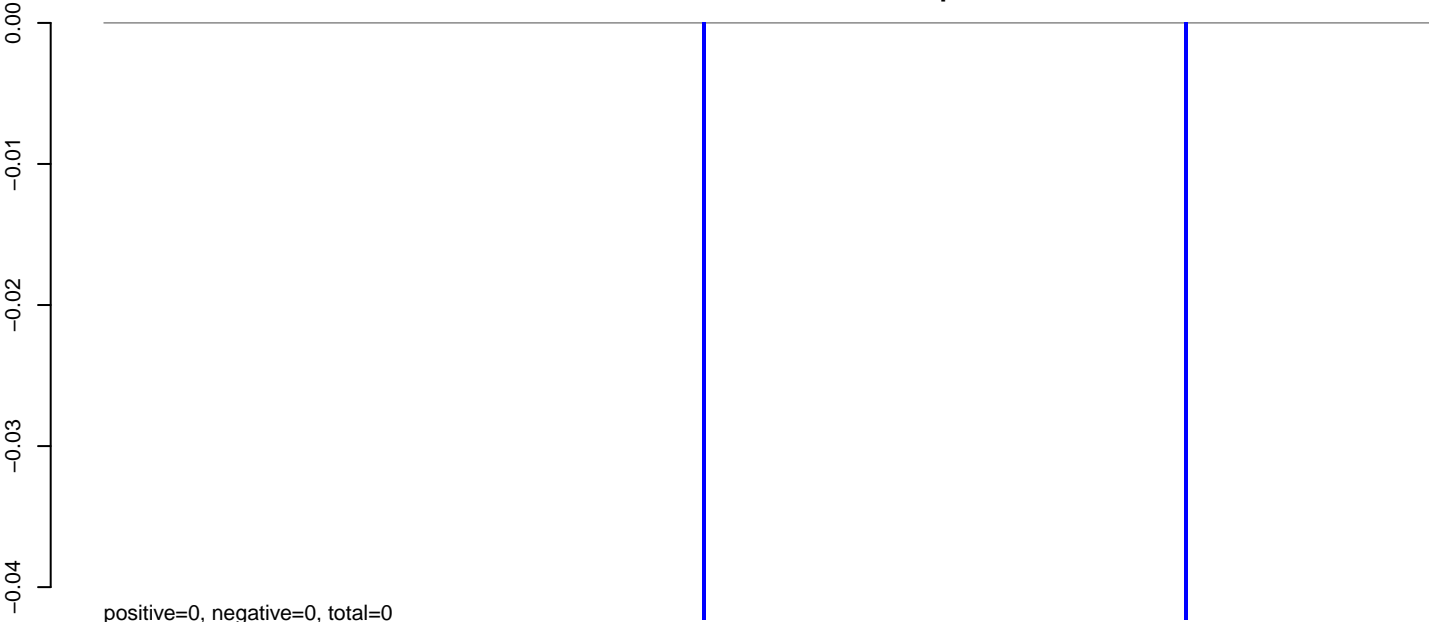
AeAlbo_C636_cells.24_35.rep



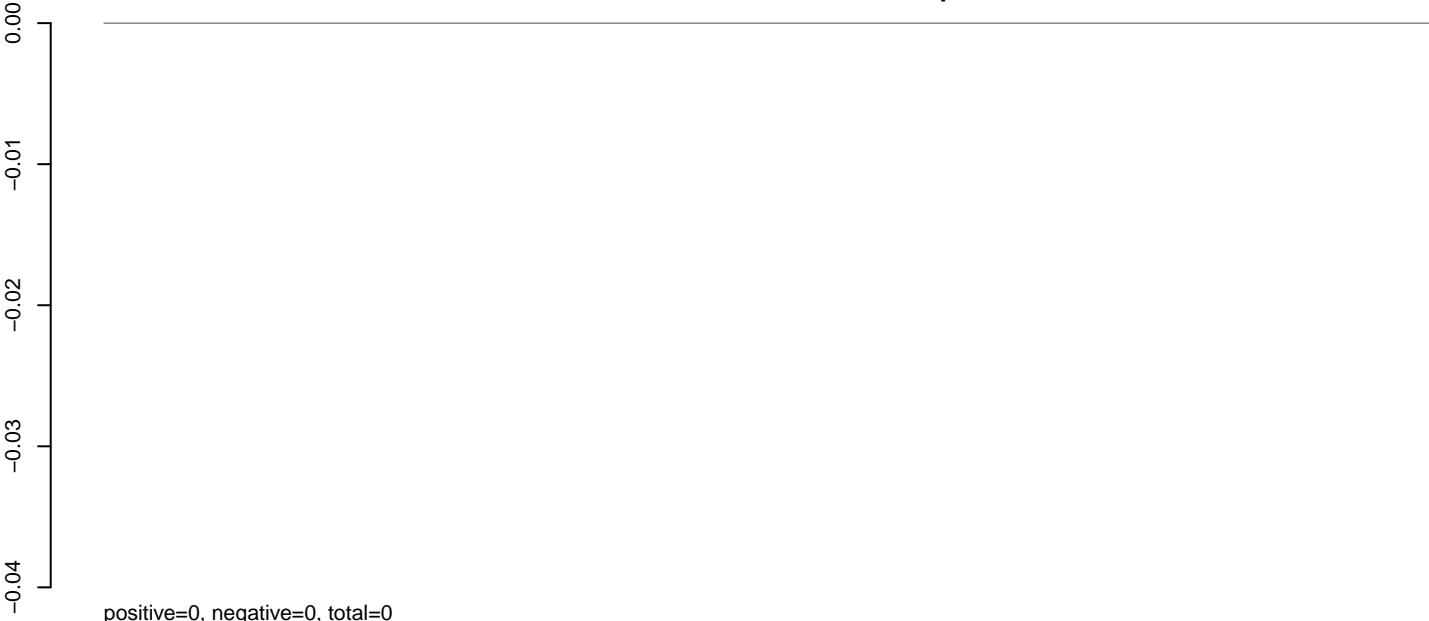
AeAlbo_C636_cells.rep



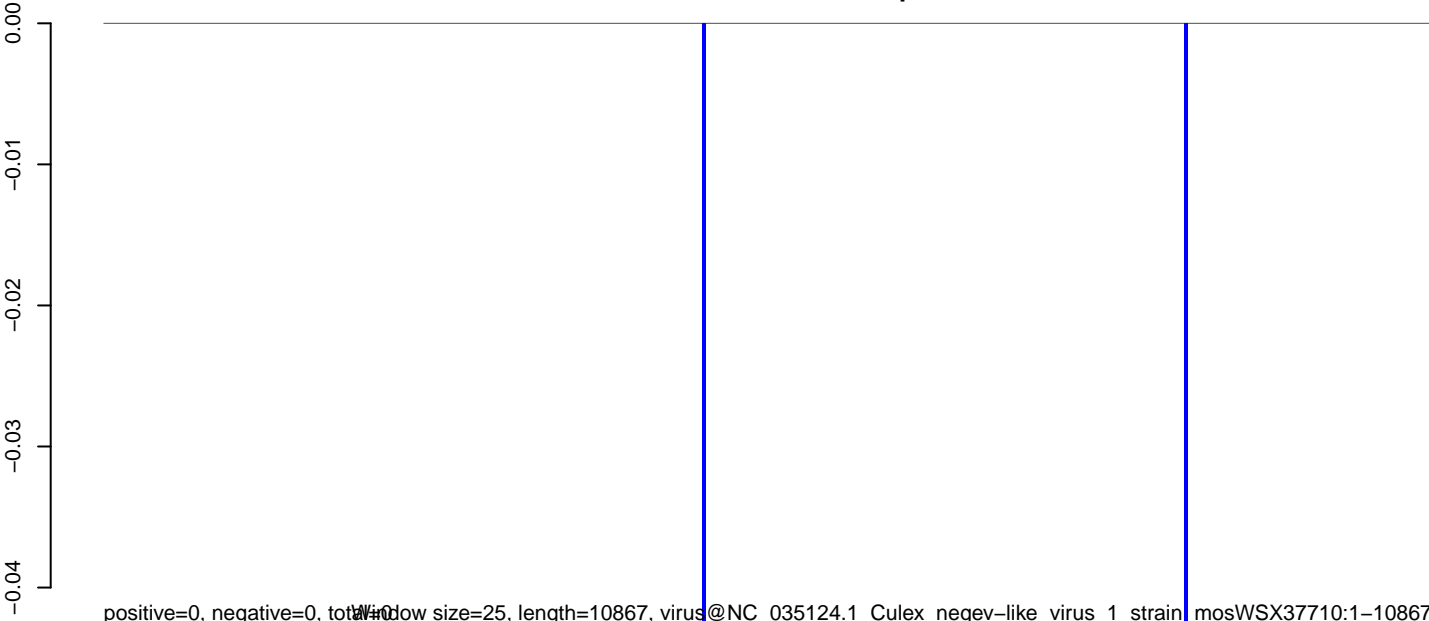
AeAlbo_C636_cells.18_23.rep



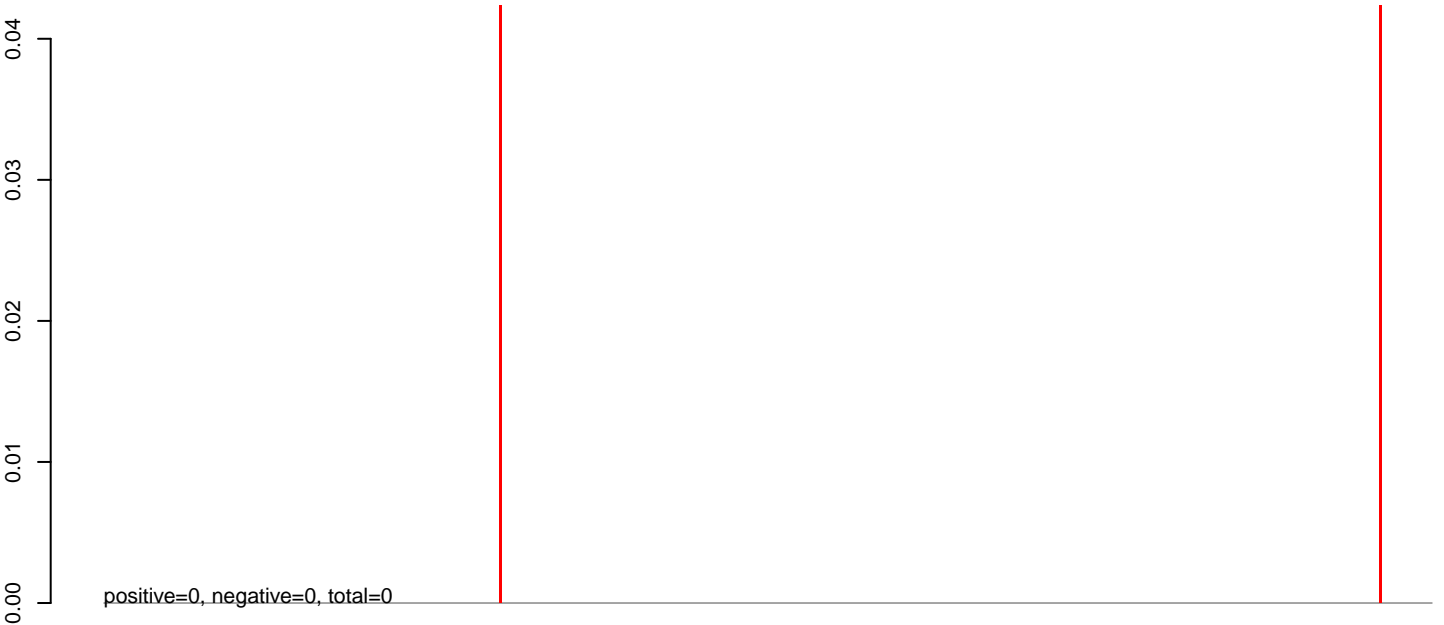
AeAlbo_C636_cells.24_35.rep



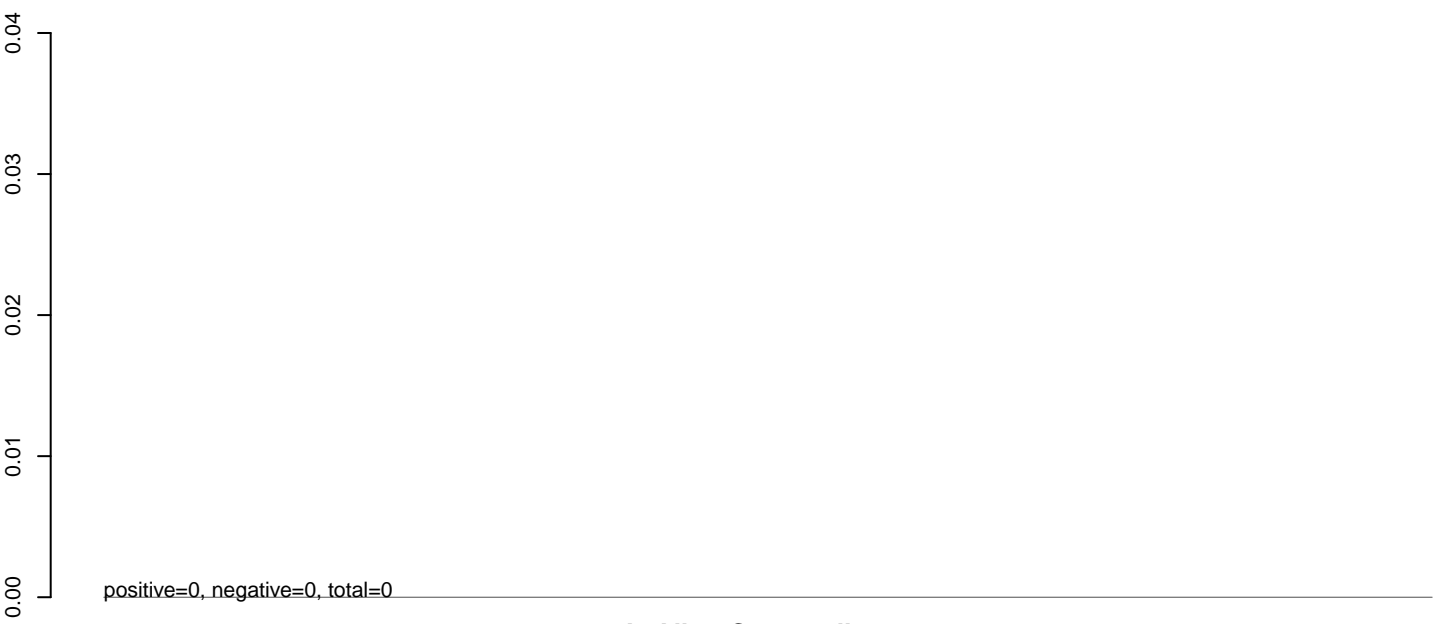
AeAlbo_C636_cells.rep



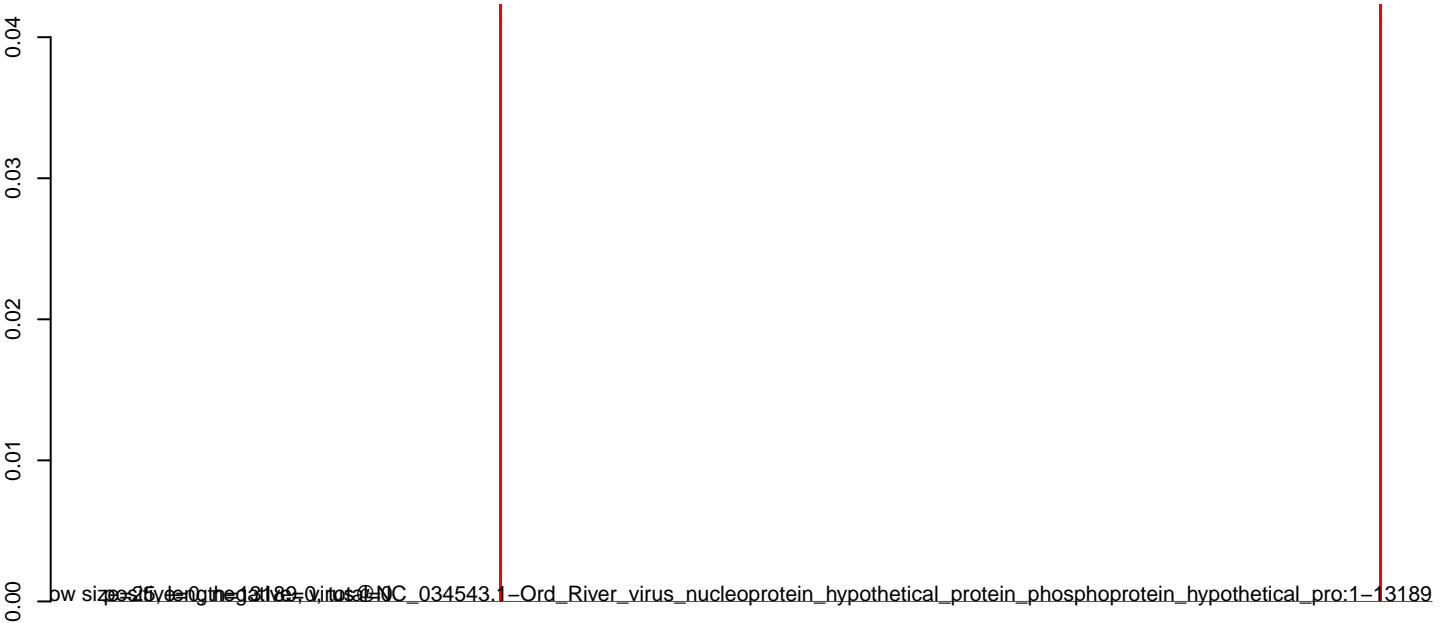
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

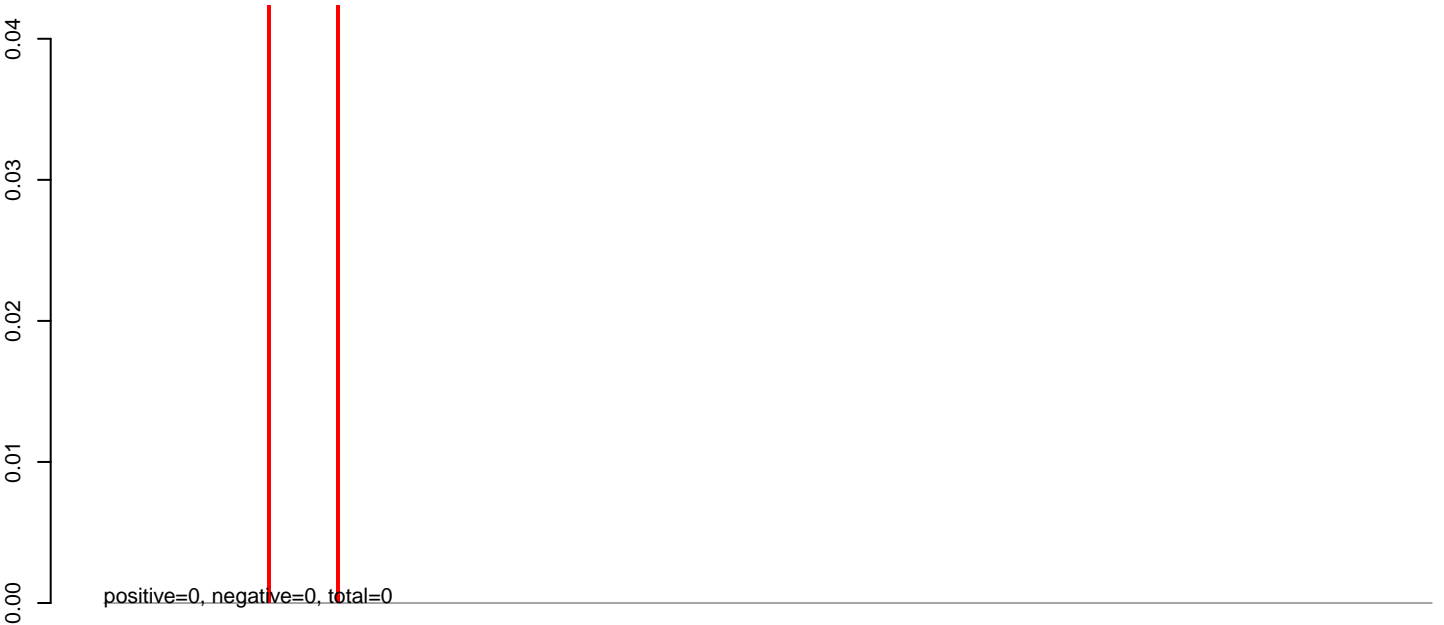


AeAlbo_C636_cells.rep

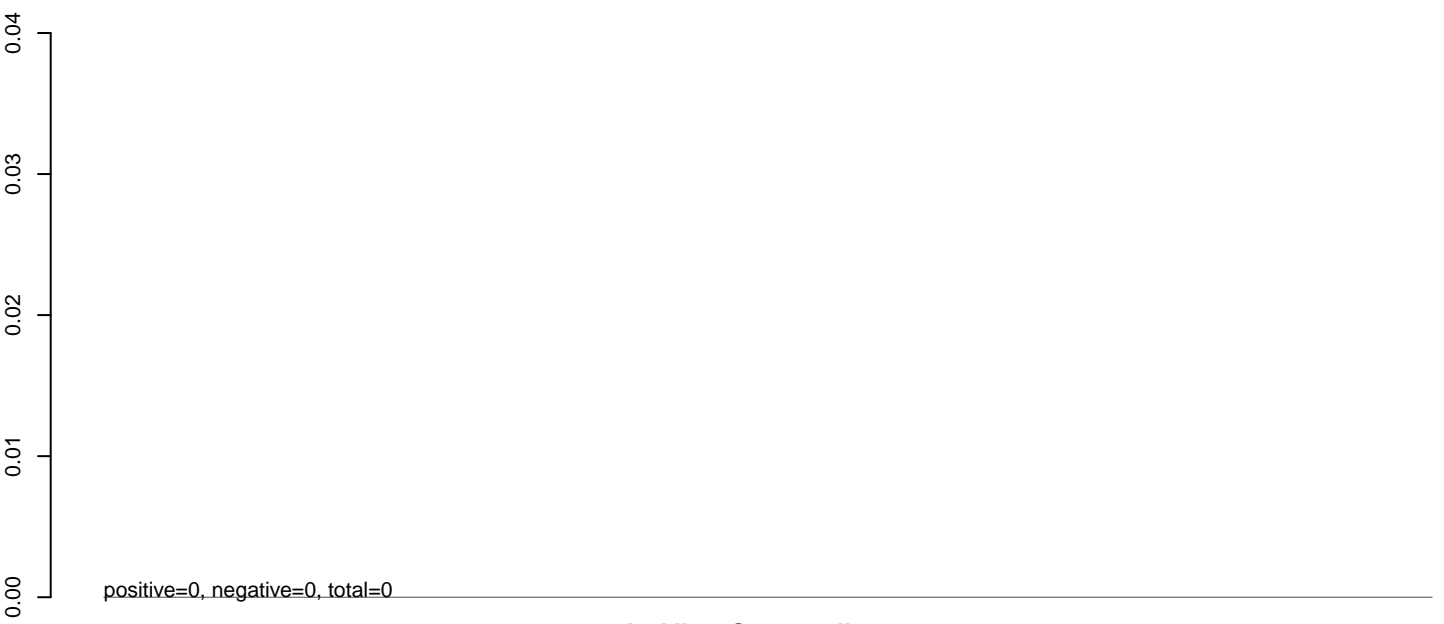


NC_034543.1-Ord_River_virus_nucleoprotein_hypothetical_protein_phosphoprotein_hypothetical_pro:1-13189

AeAlbo_C636_cells.18_23.rep



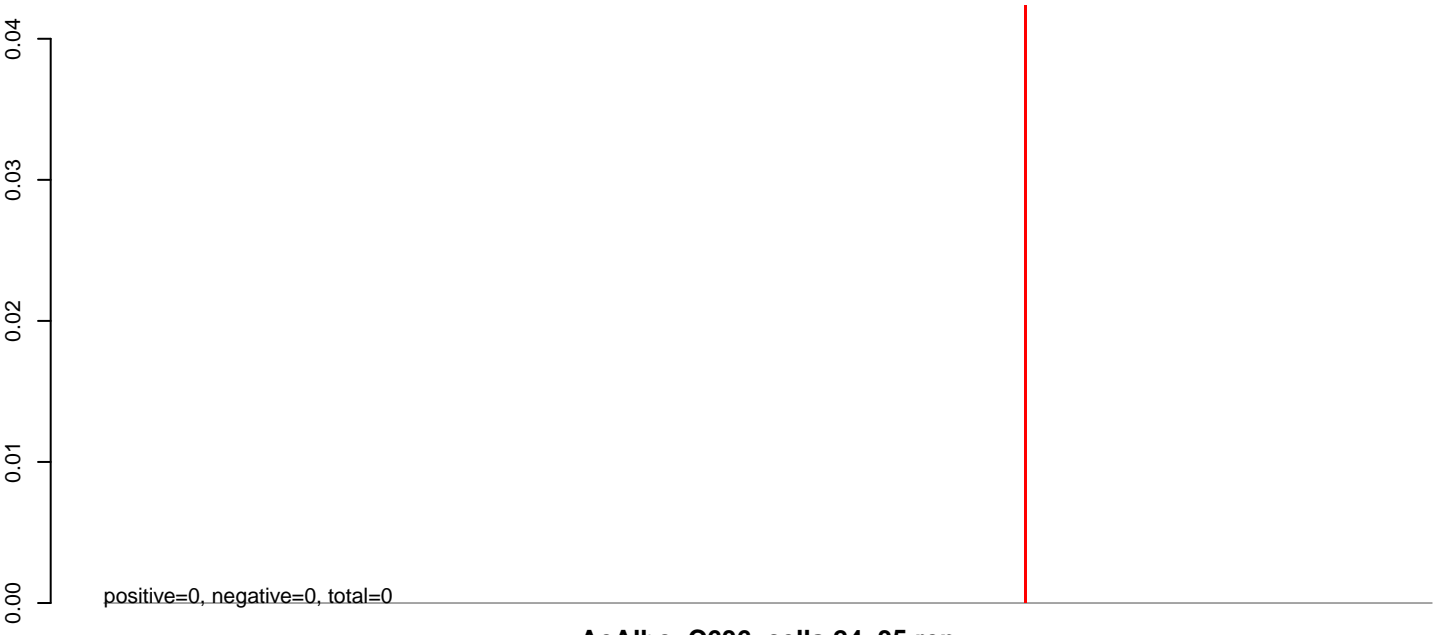
AeAlbo_C636_cells.24_35.rep



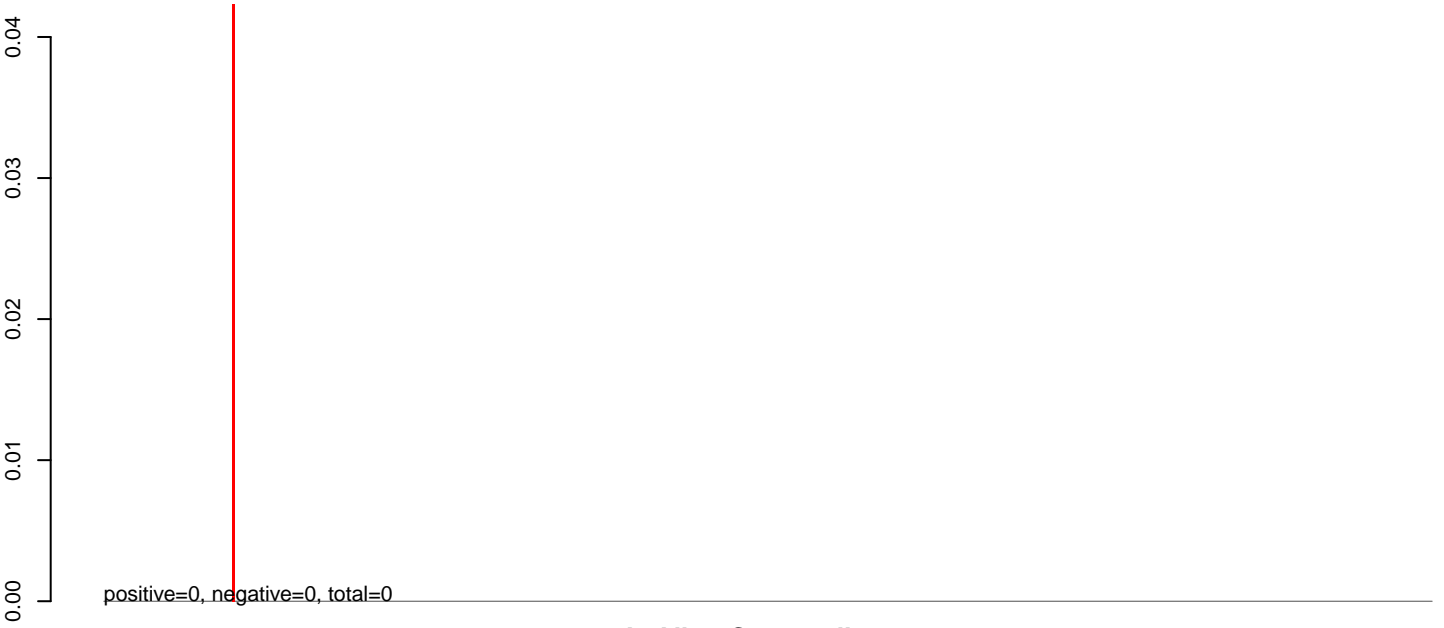
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



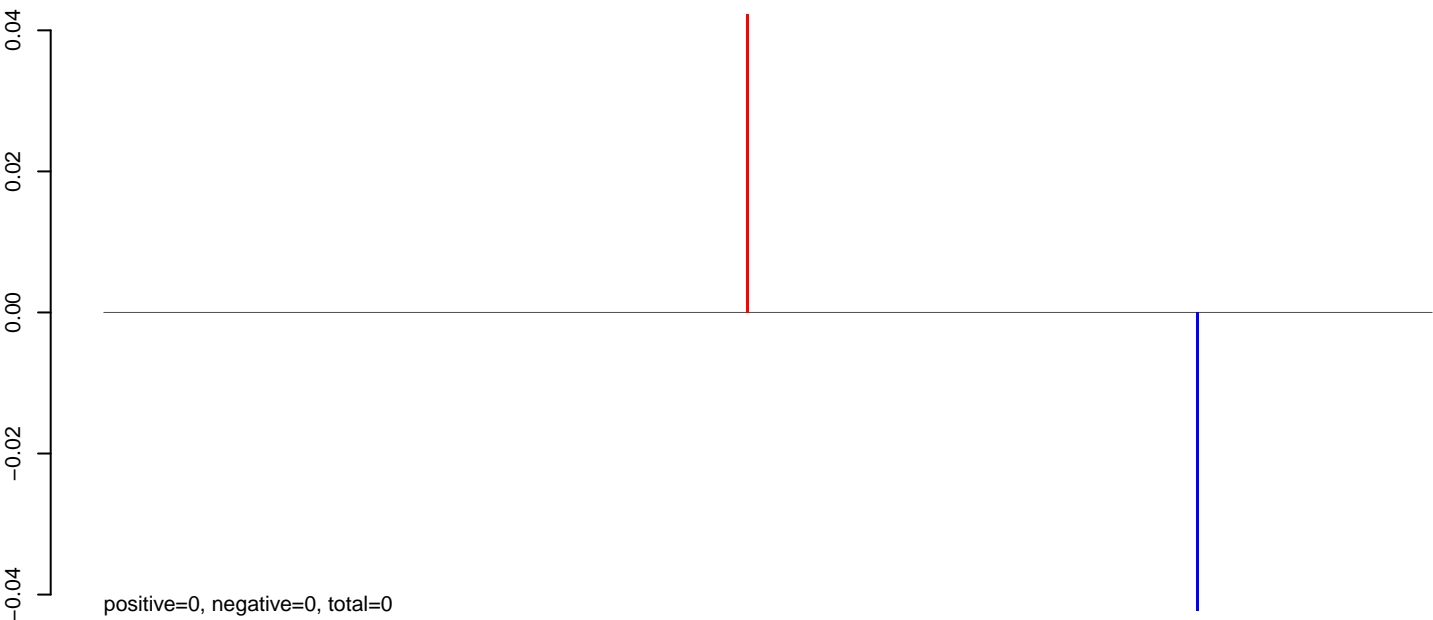
AeAlbo_C636_cells.24_35.rep



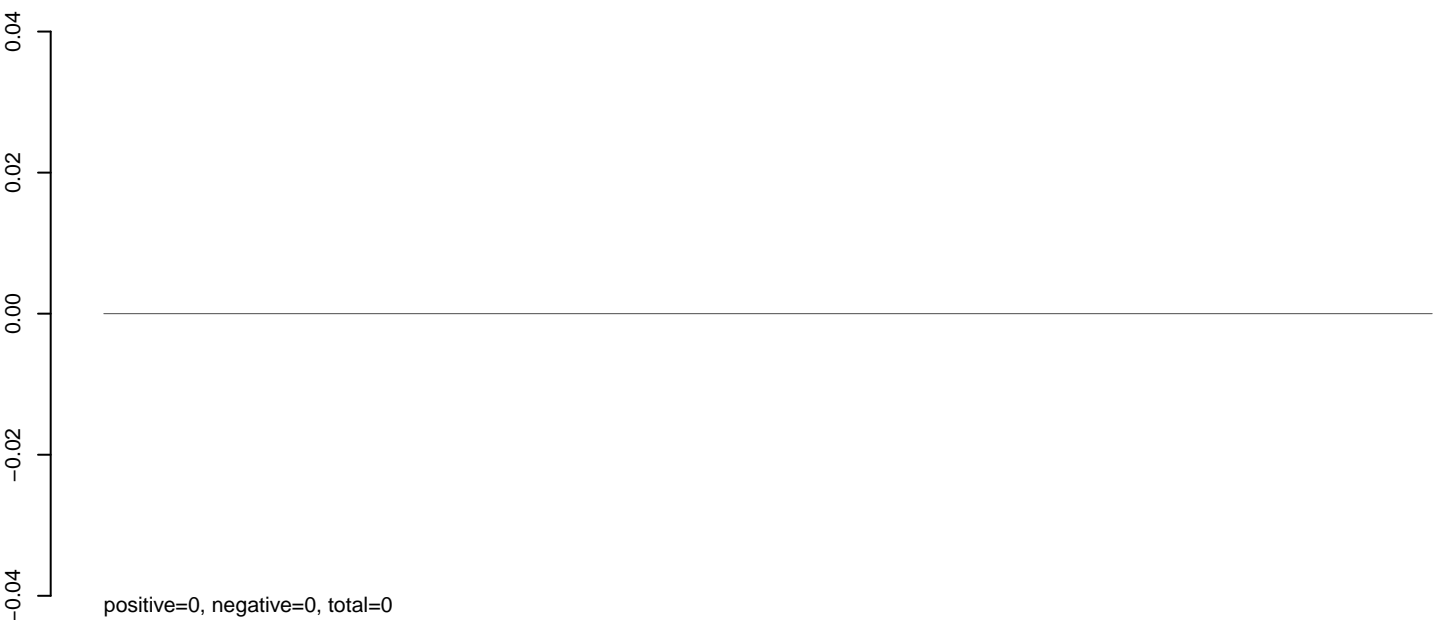
AeAlbo_C636_cells.rep



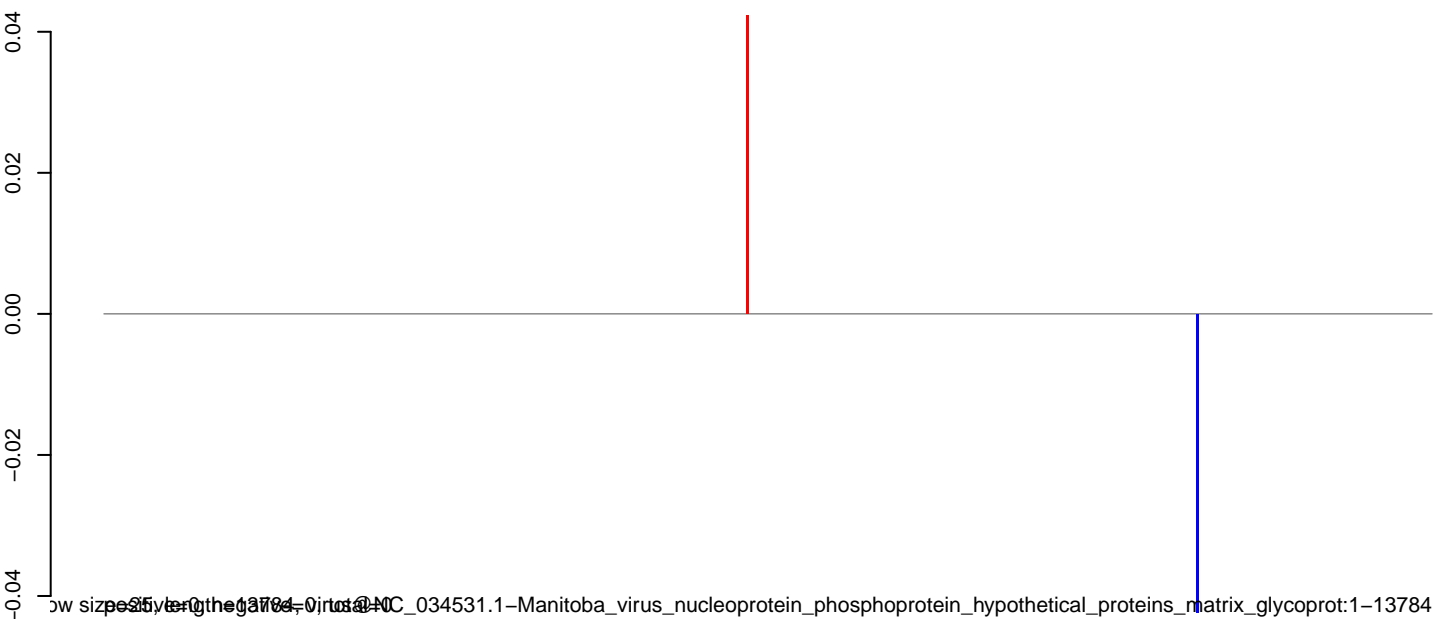
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



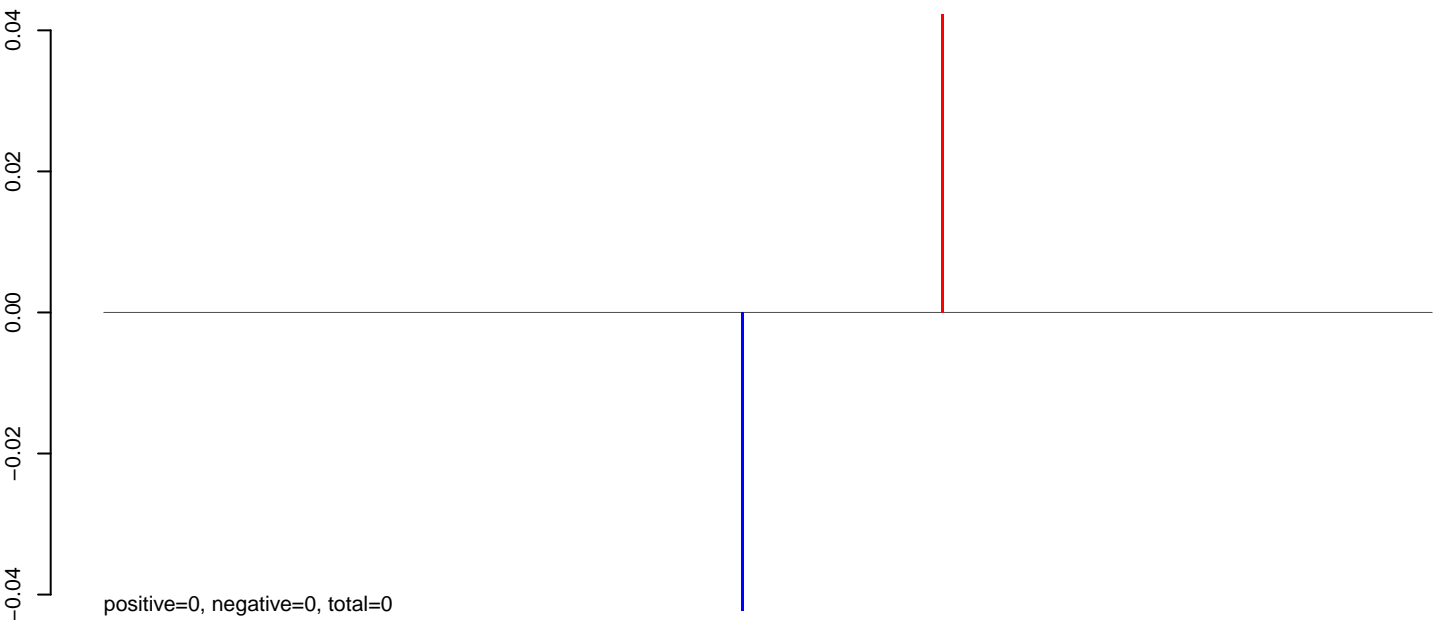
AeAlbo_C636_cells.rep



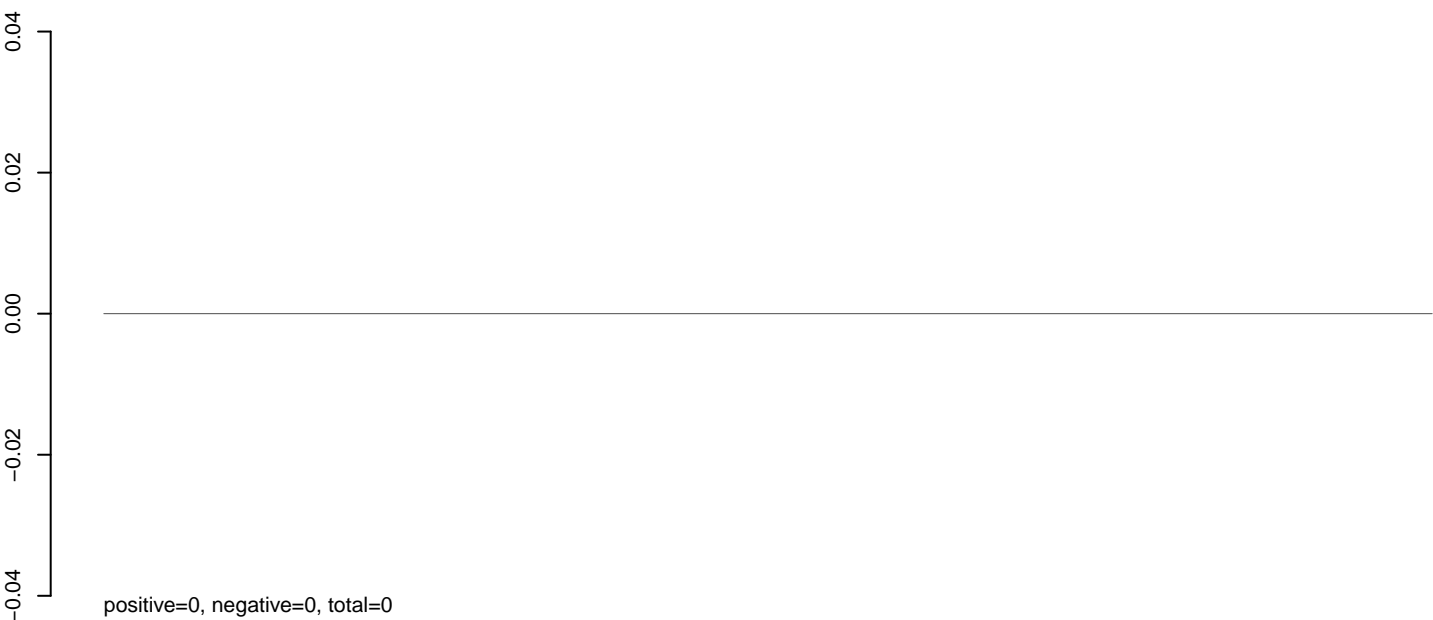
bow size=0, length=1784, int=0, NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

0 2000 4000 6000 8000 10000 12000 14000

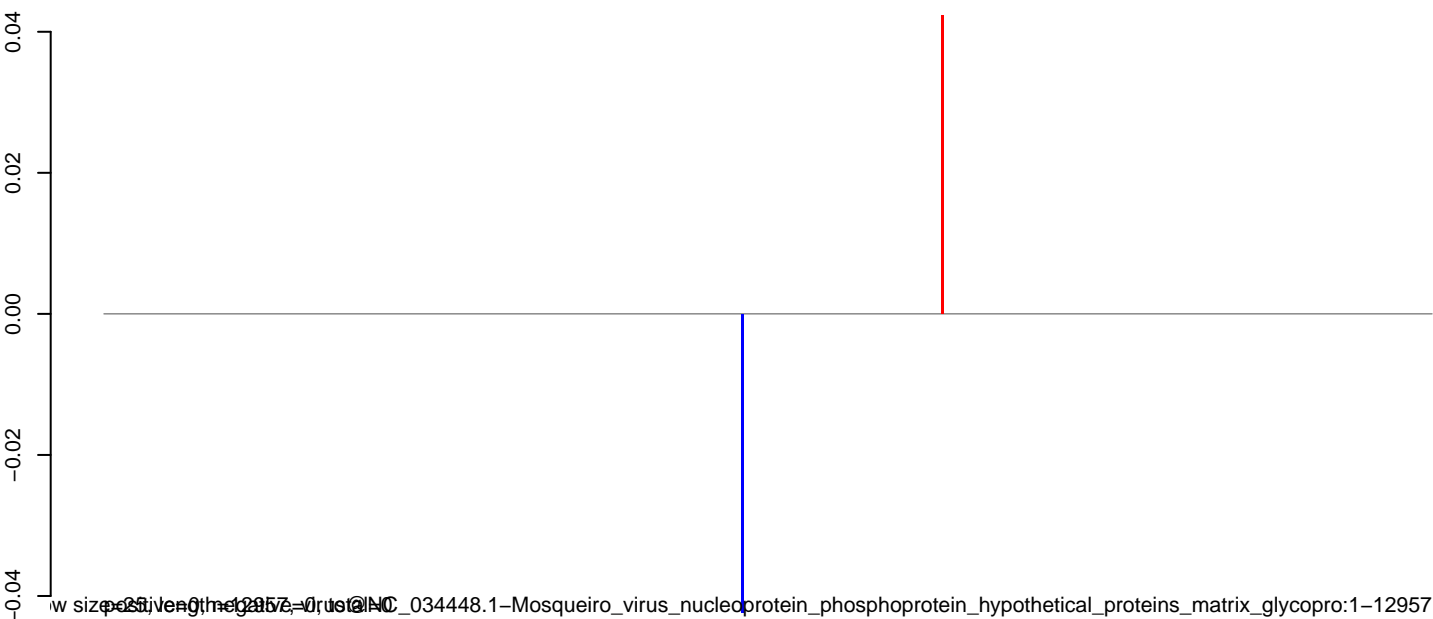
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



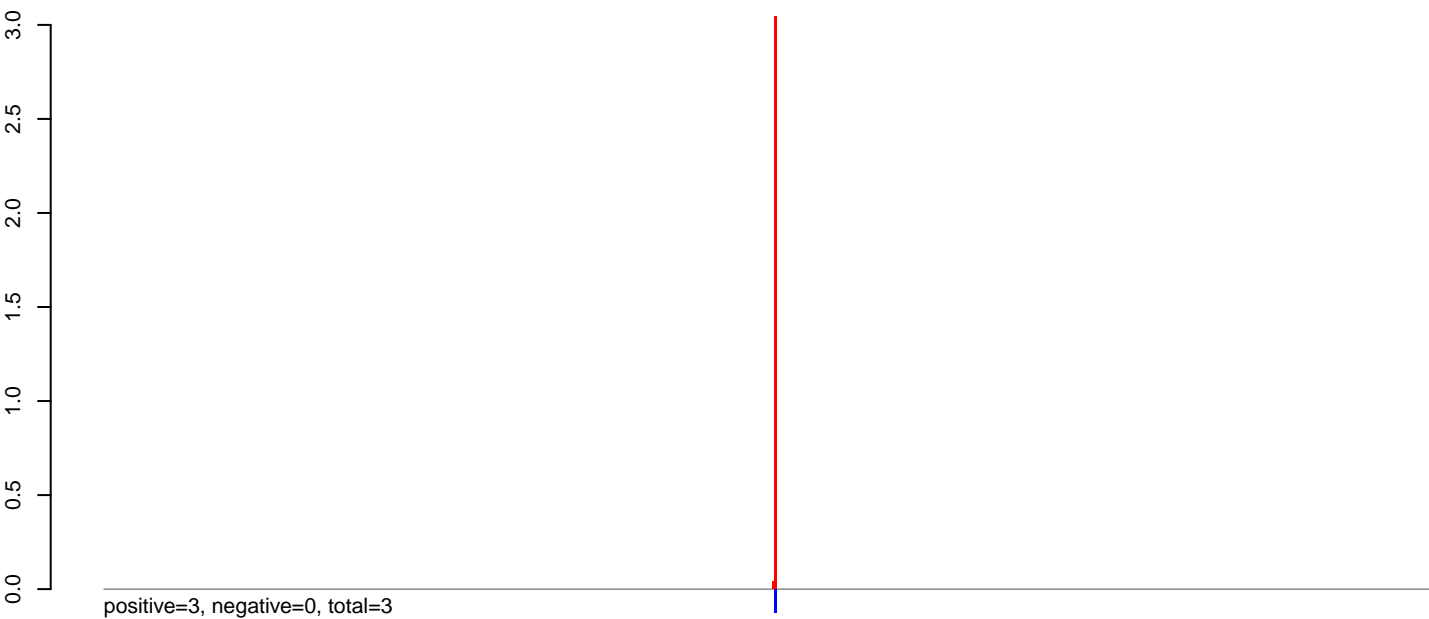
AeAlbo_C636_cells.rep



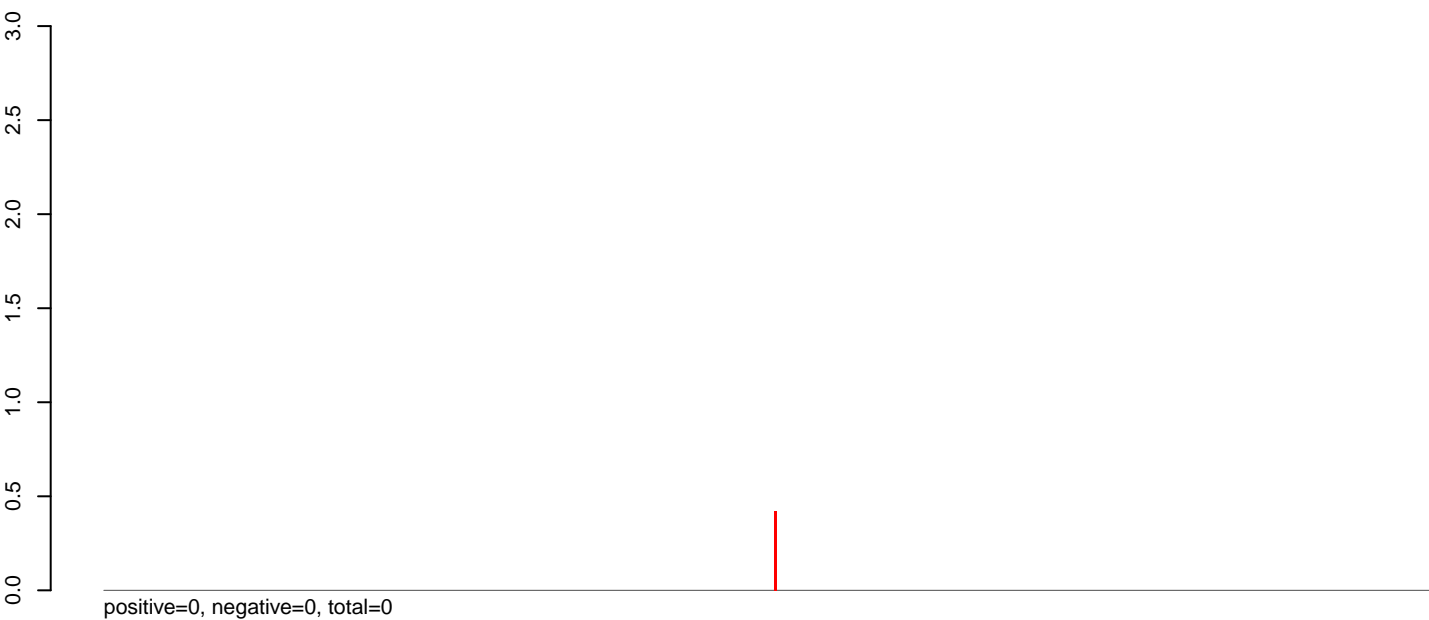
w size=25, length=12957, use@NC_034448.1-Mosqueiro_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-12957

0 2000 4000 6000 8000 10000 12000

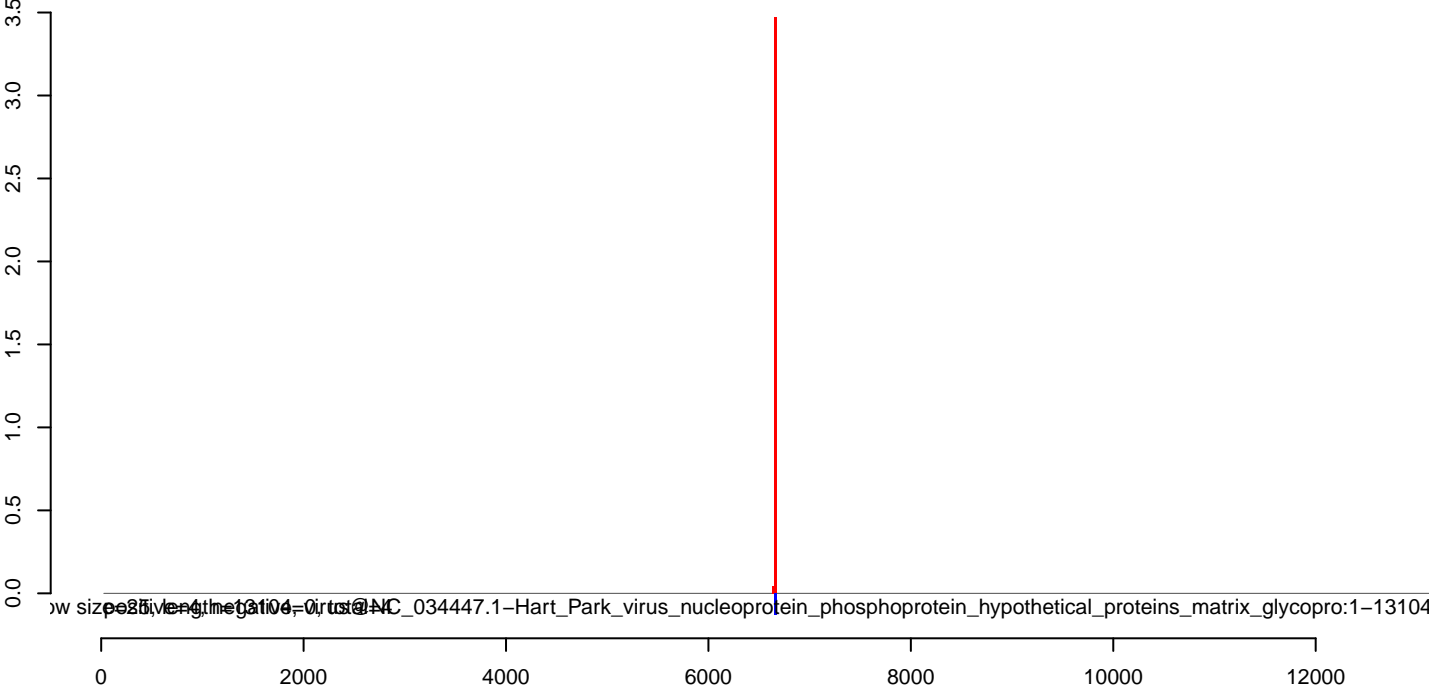
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

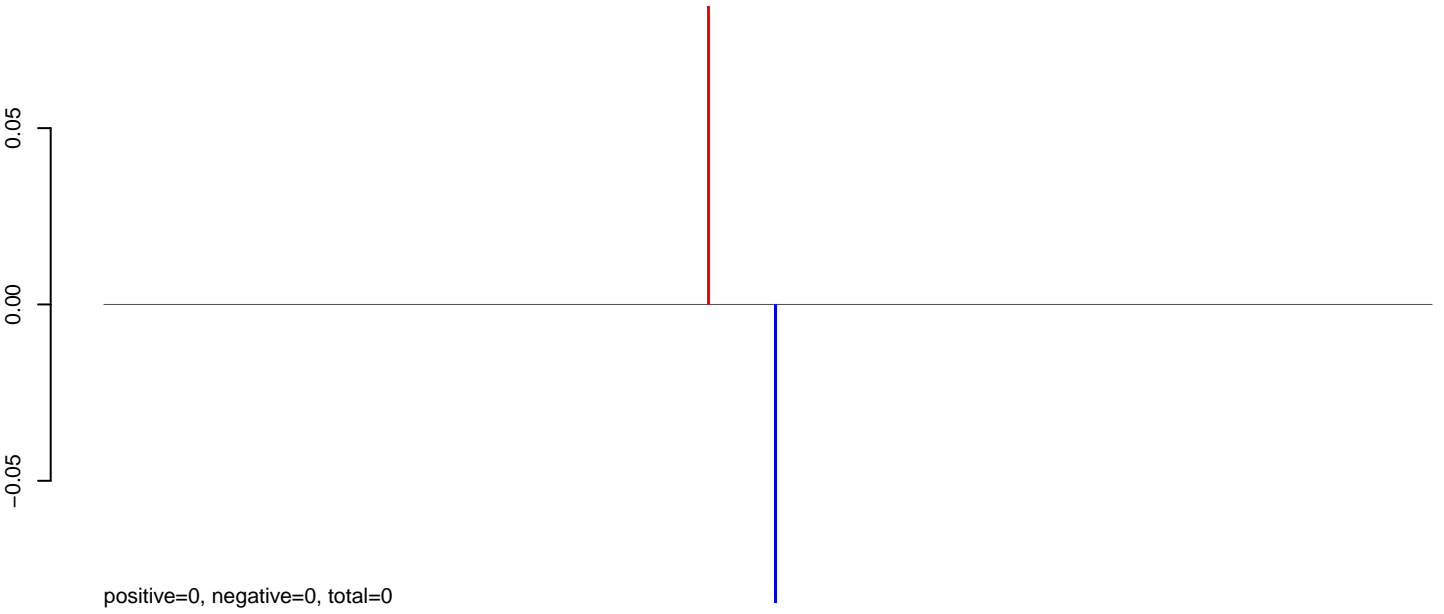


AeAlbo_C636_cells.rep

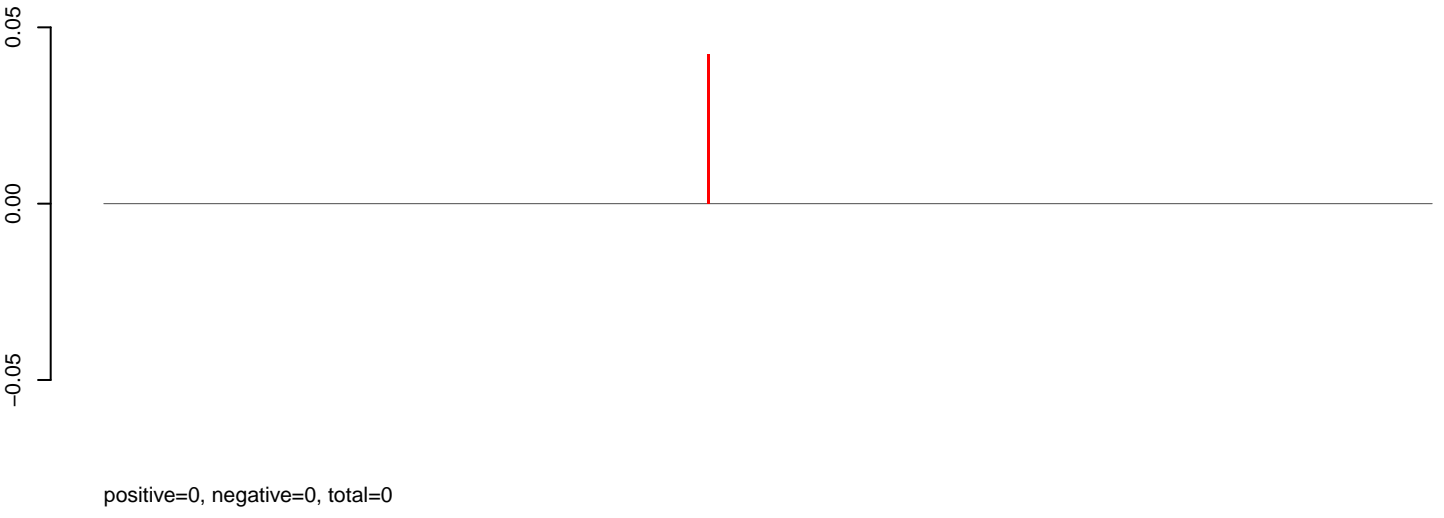


ww size positive=1, negative=0, total=1
NC_034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

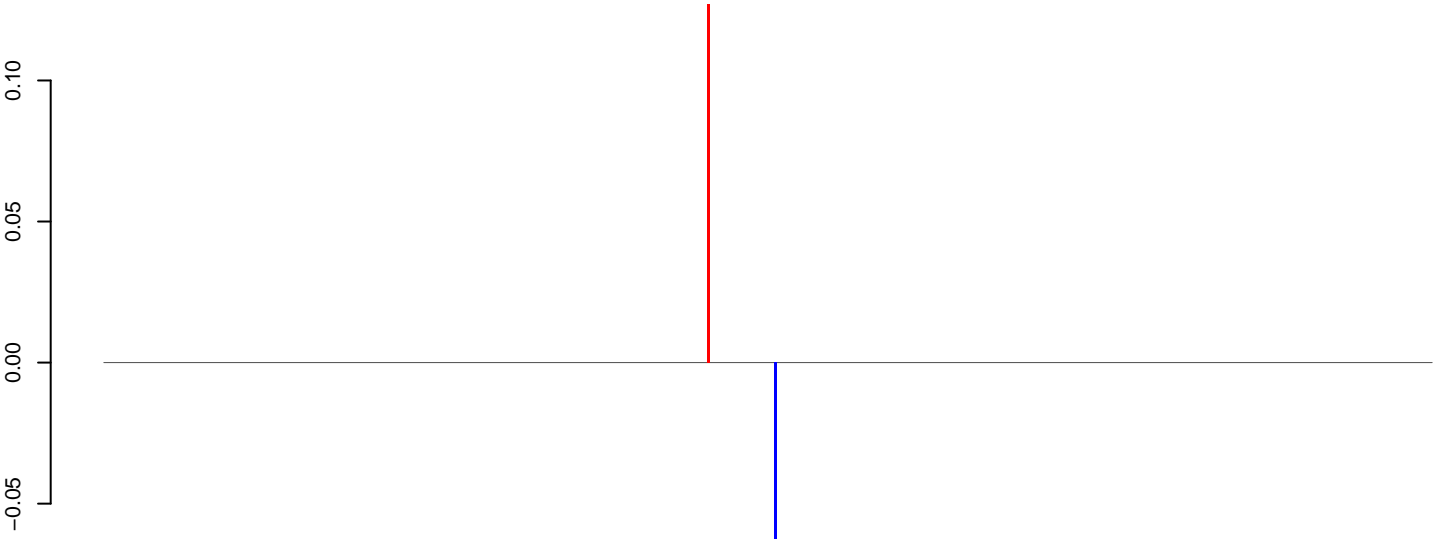
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



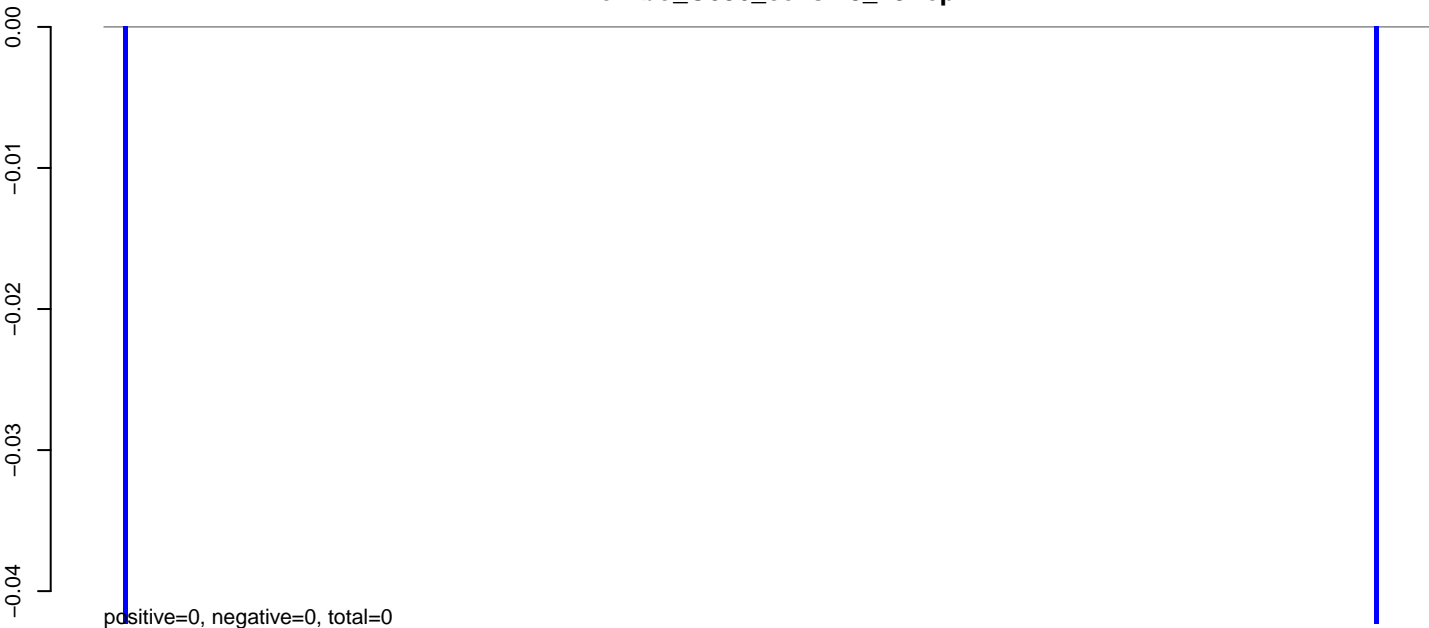
AeAlbo_C636_cells.rep



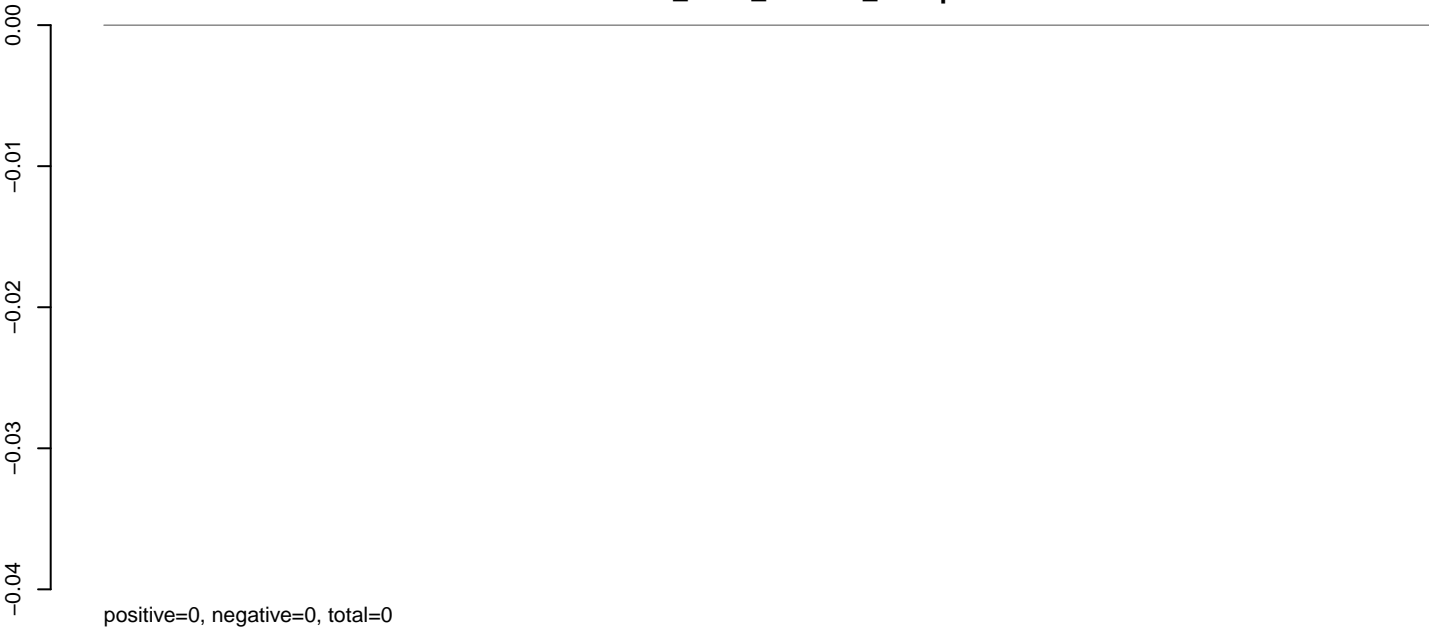
www.ebi.ac.uk/ena/submit/150-0-us@NC_034443.1-Le_Dantec_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_prot:1-11450

0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

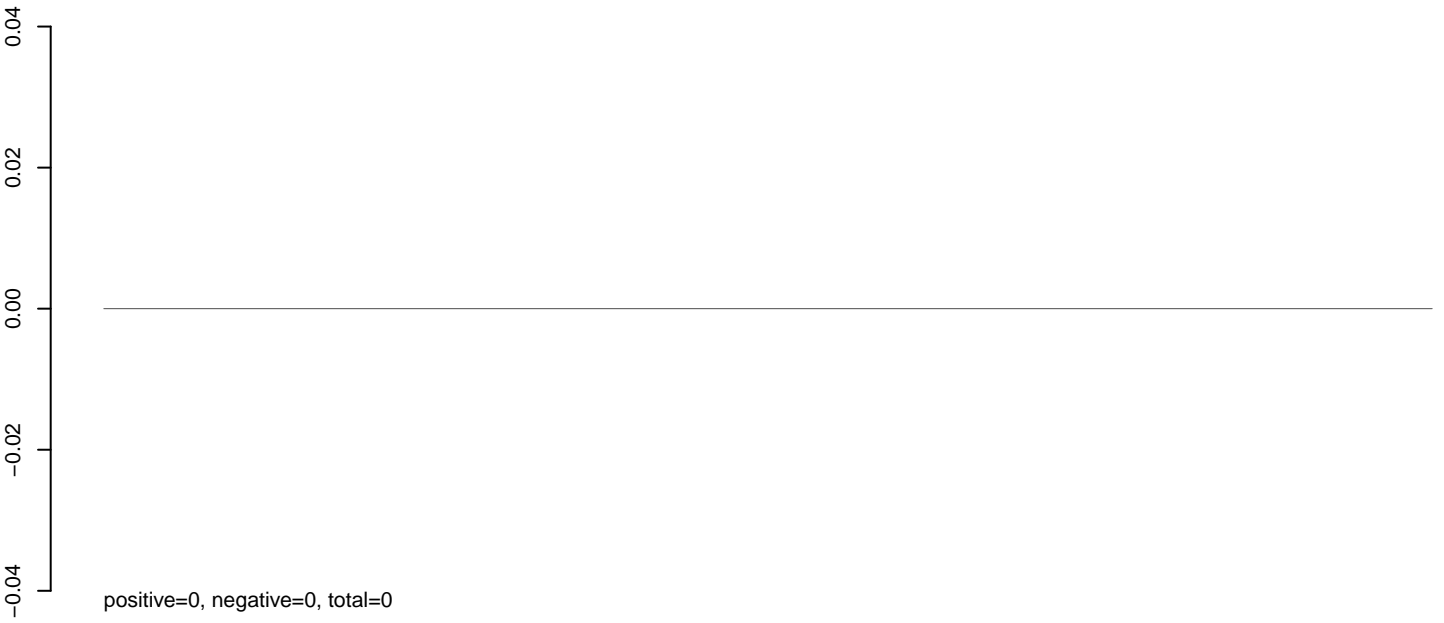


0 1000 2000 3000 4000 5000 6000 7000

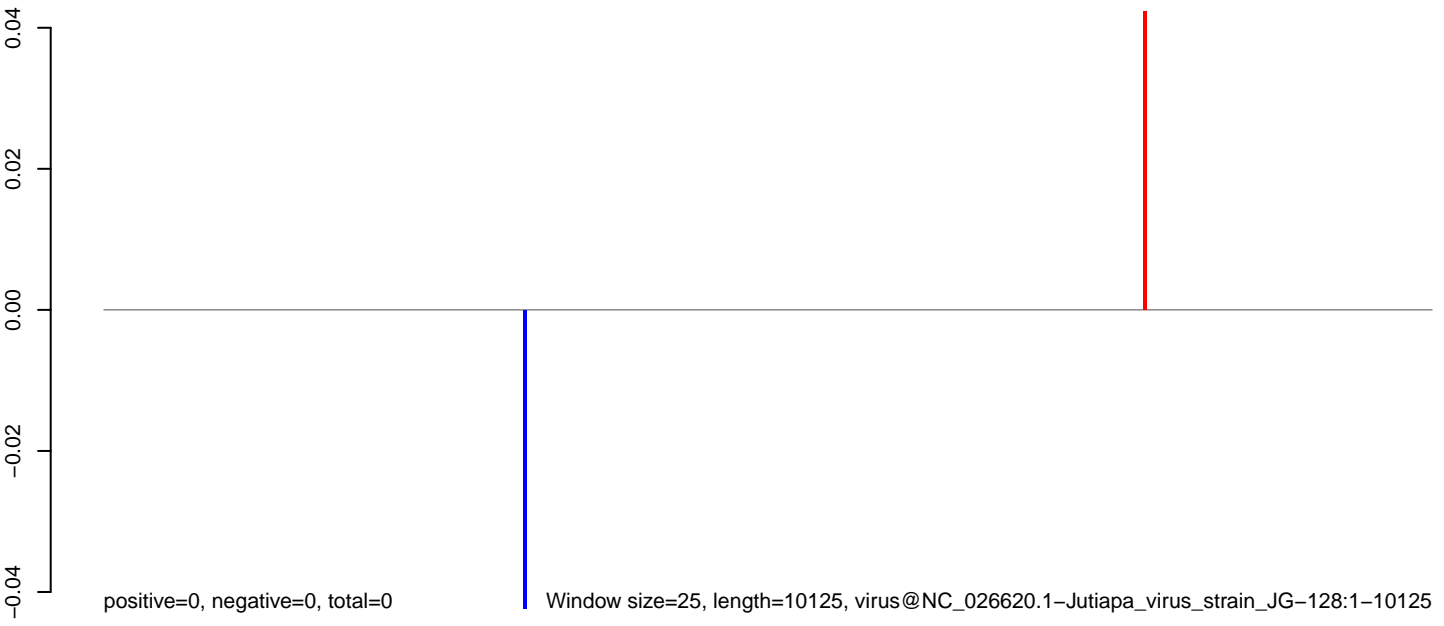
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

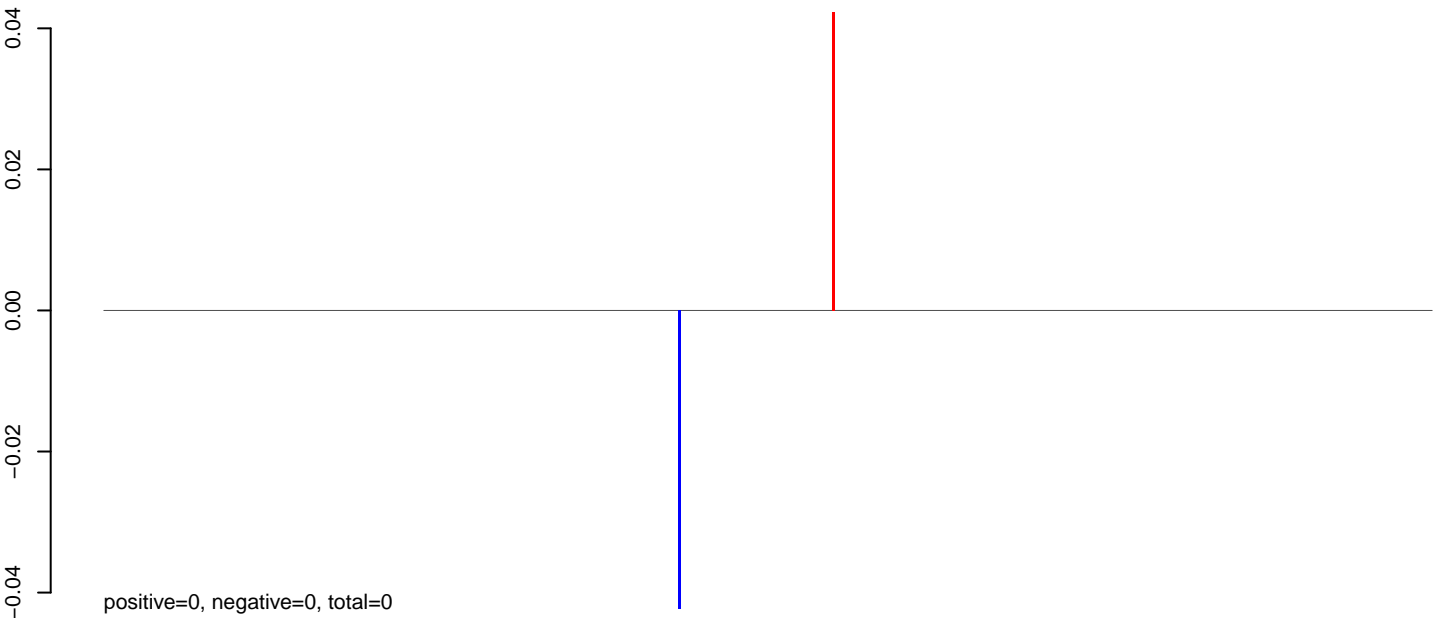


AeAlbo_C636_cells.rep

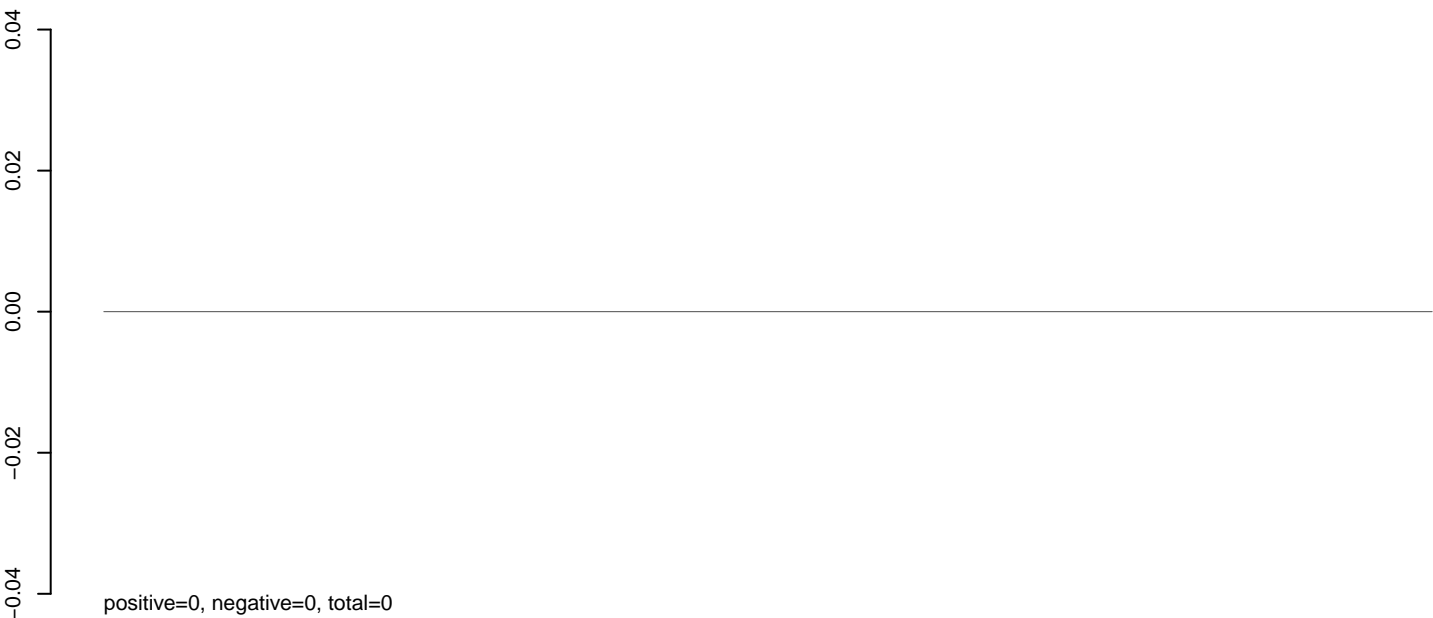


0 2000 4000 6000 8000 10000

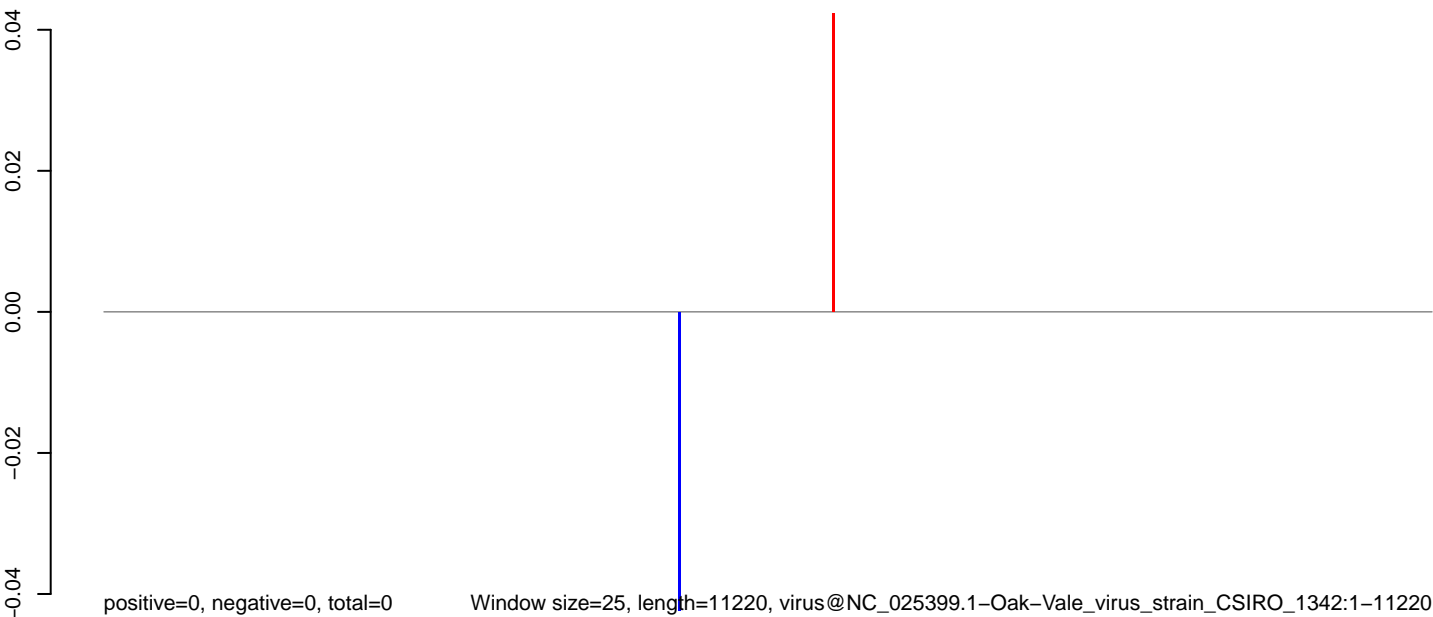
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

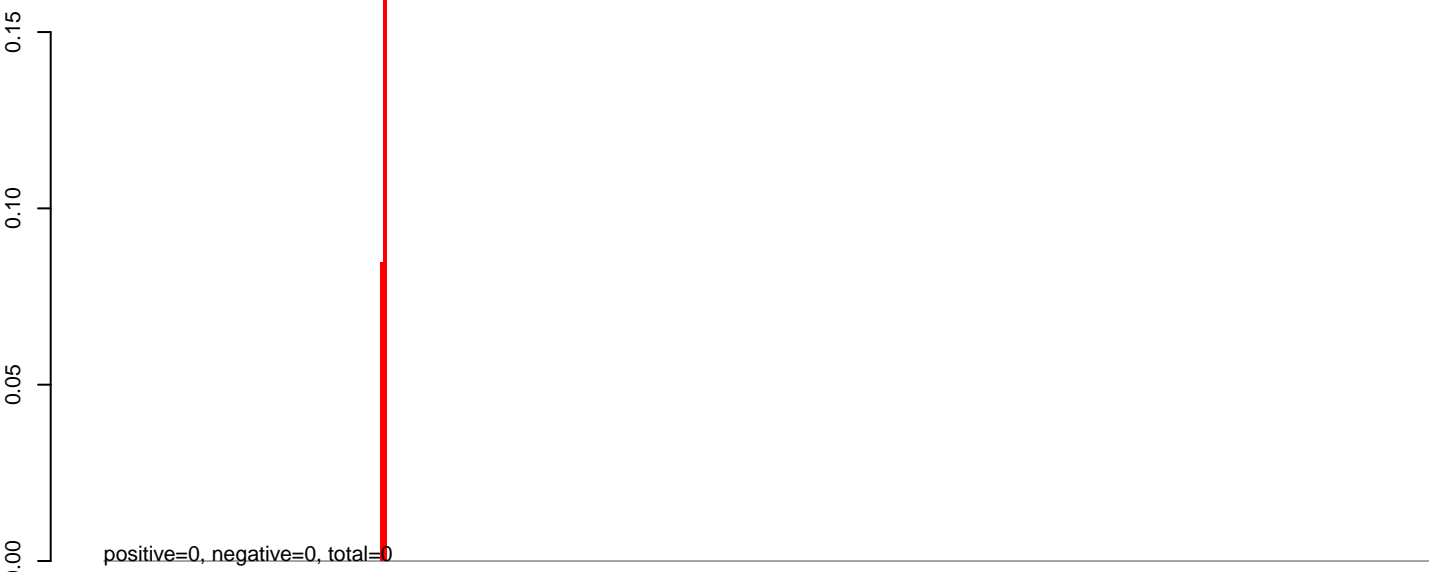


AeAlbo_C636_cells.rep

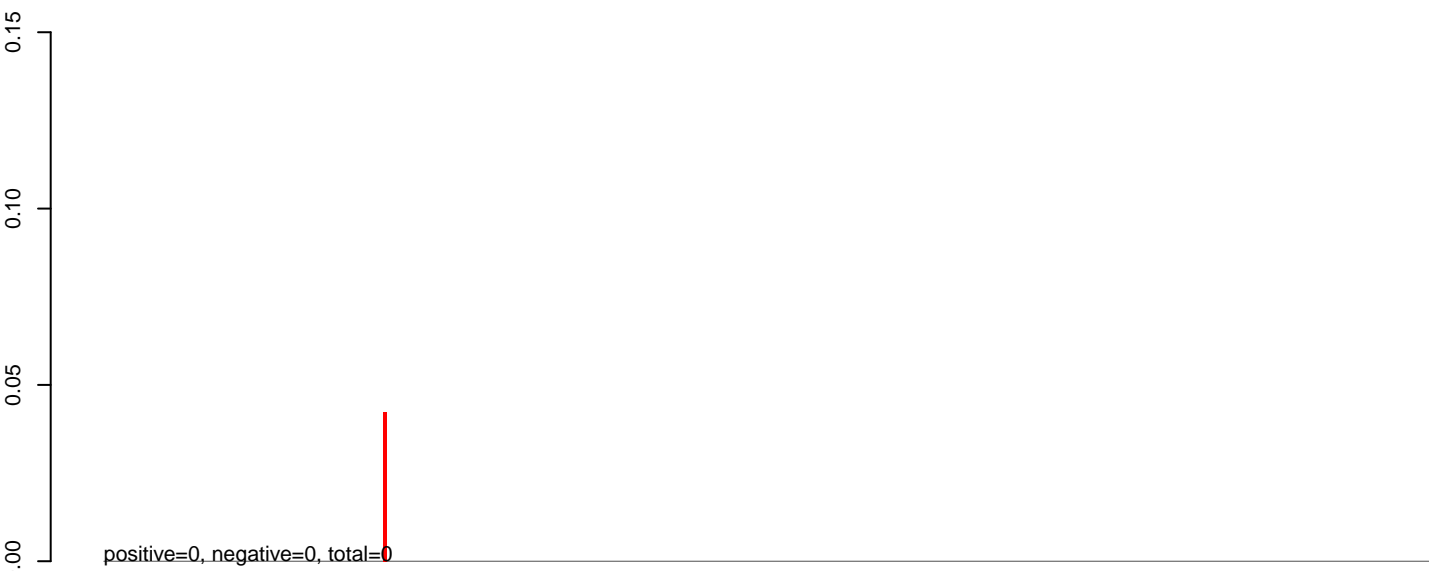


0 2000 4000 6000 8000 10000

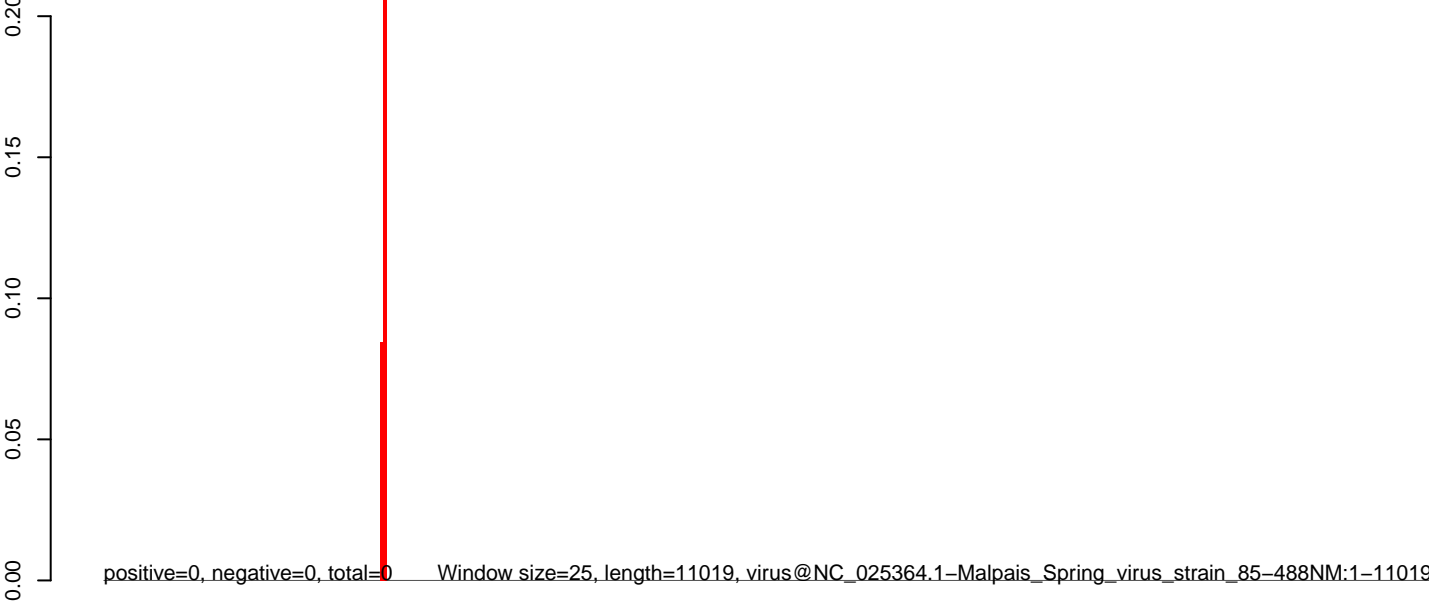
AeAlbo_C636_cells.18_23.rep



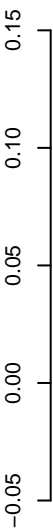
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

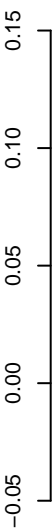


AeAlbo_C636_cells.18_23.rep



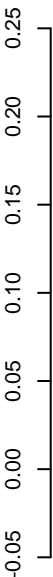
positive=0, negative=0, total=1

AeAlbo_C636_cells.24_35.rep

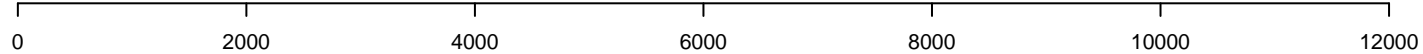


positive=0, negative=0, total=0

AeAlbo_C636_cells.rep



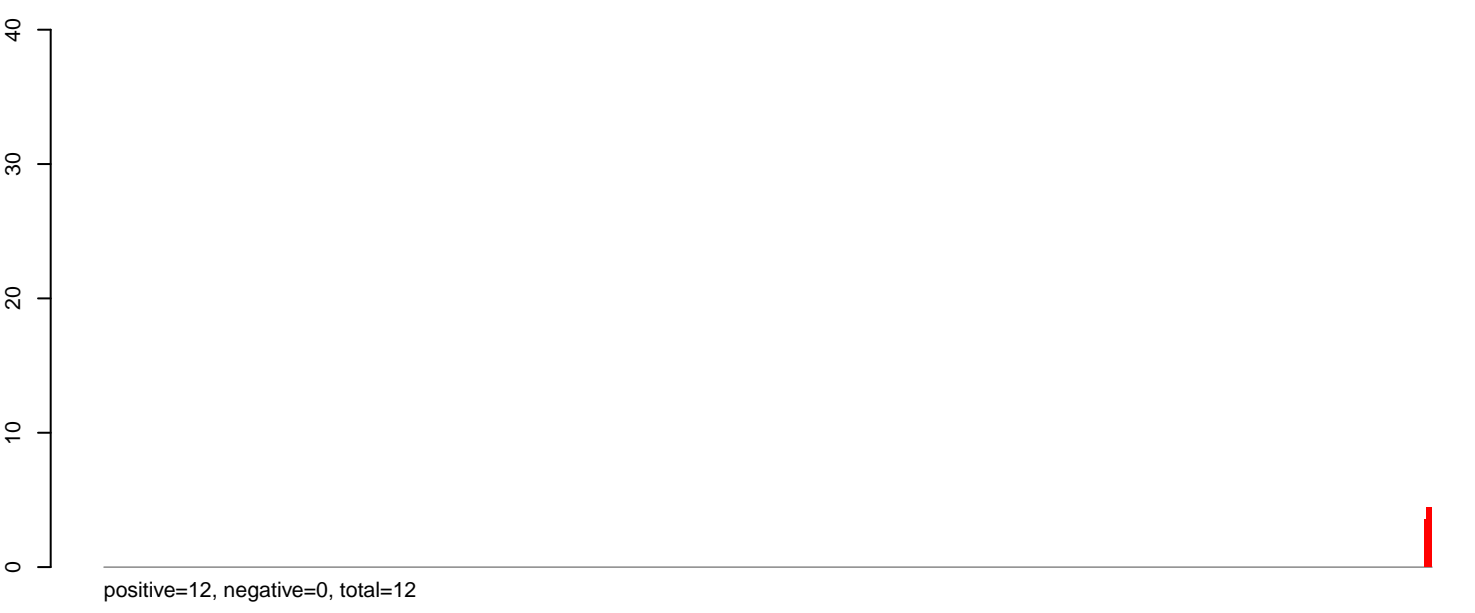
positive=1, negative=25, total=26, length=11624, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624



AeAlbo_C636_cells.18_23.rep



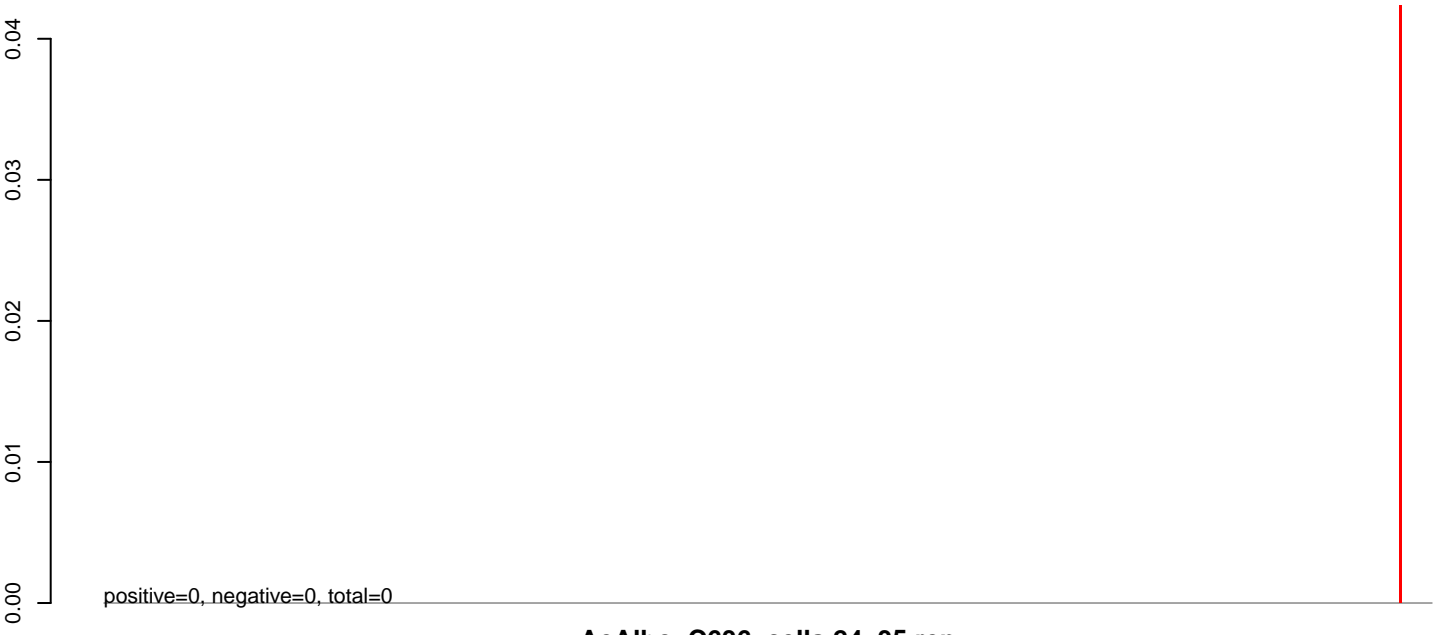
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



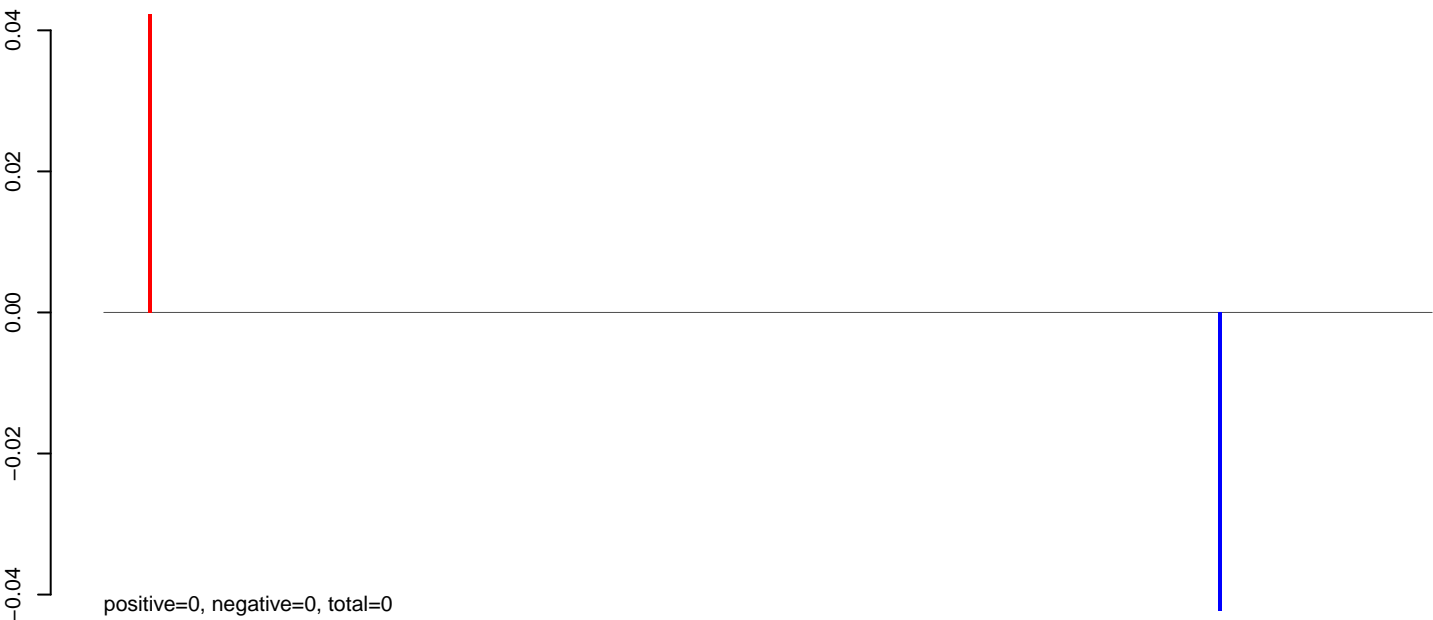
AeAlbo_C636_cells.24_35.rep



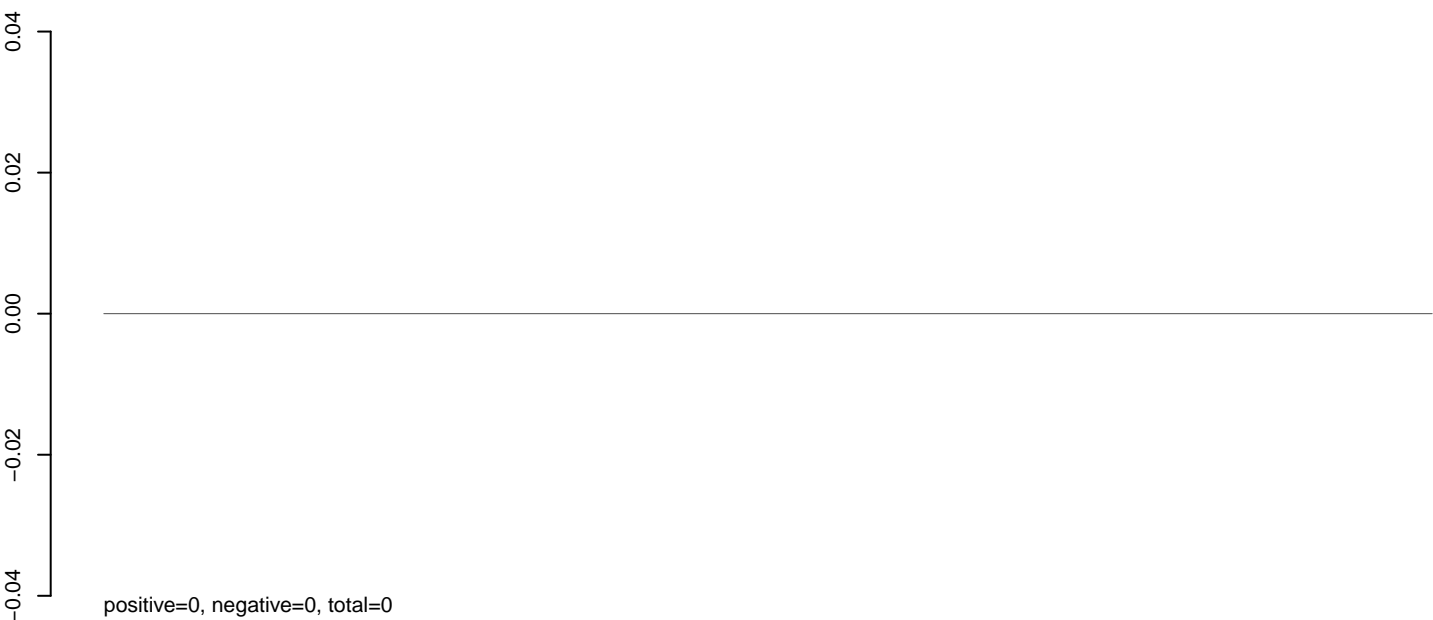
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

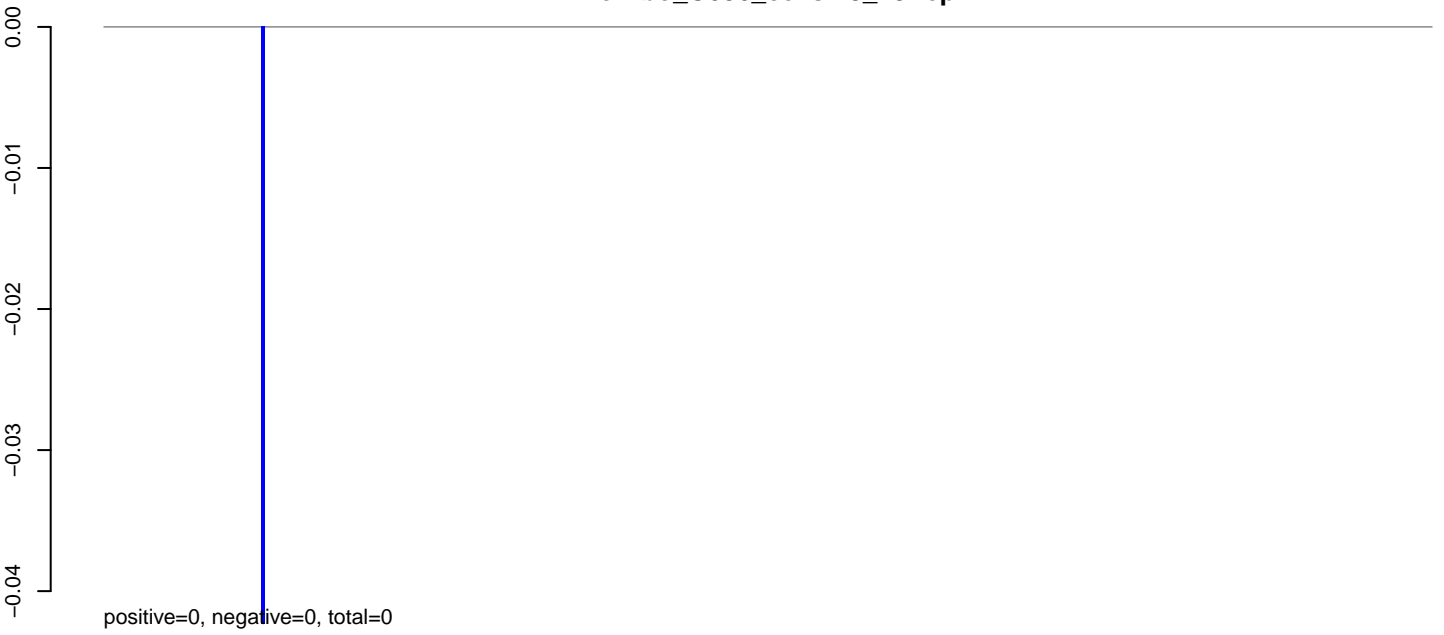


AeAlbo_C636_cells.rep

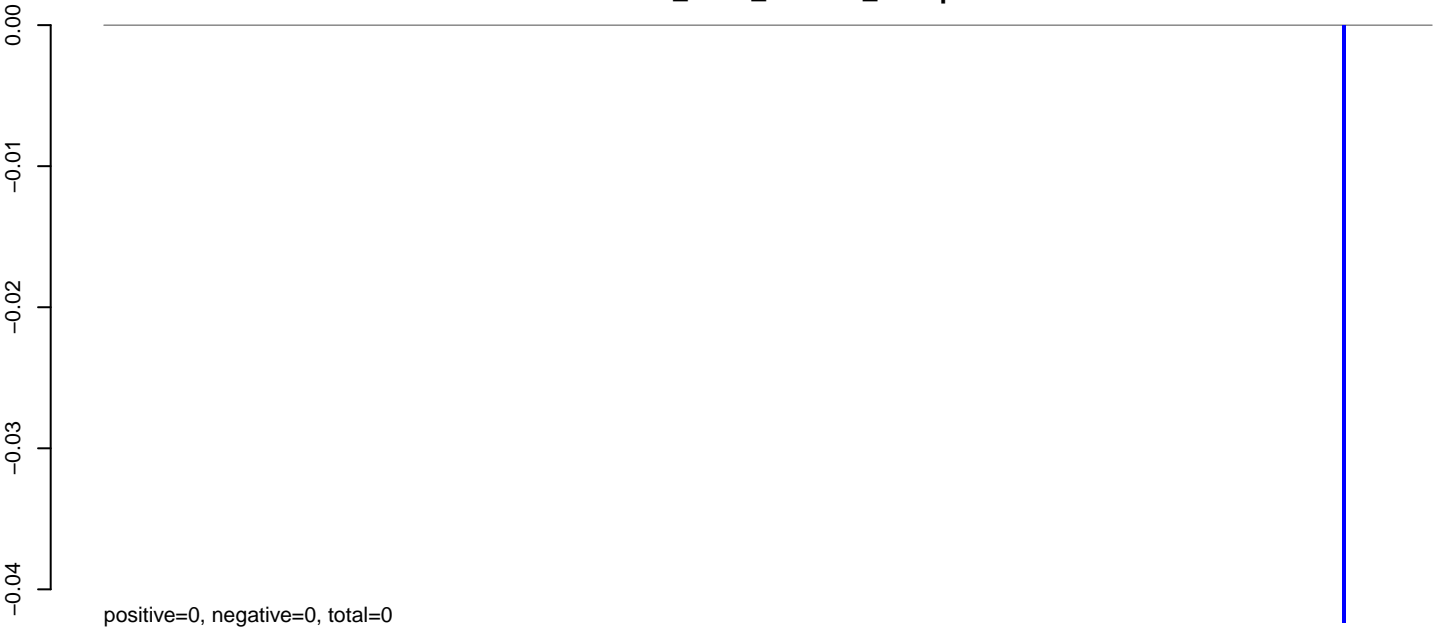


0 2000 4000 6000 8000 10000

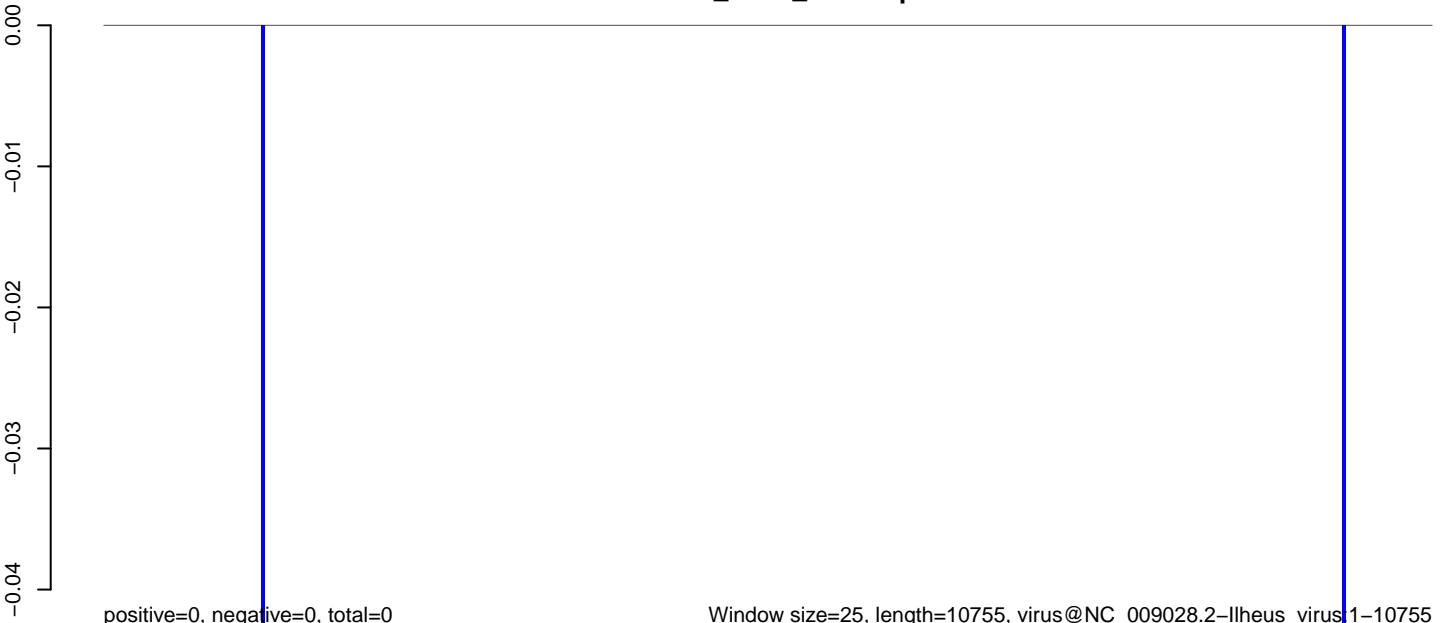
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



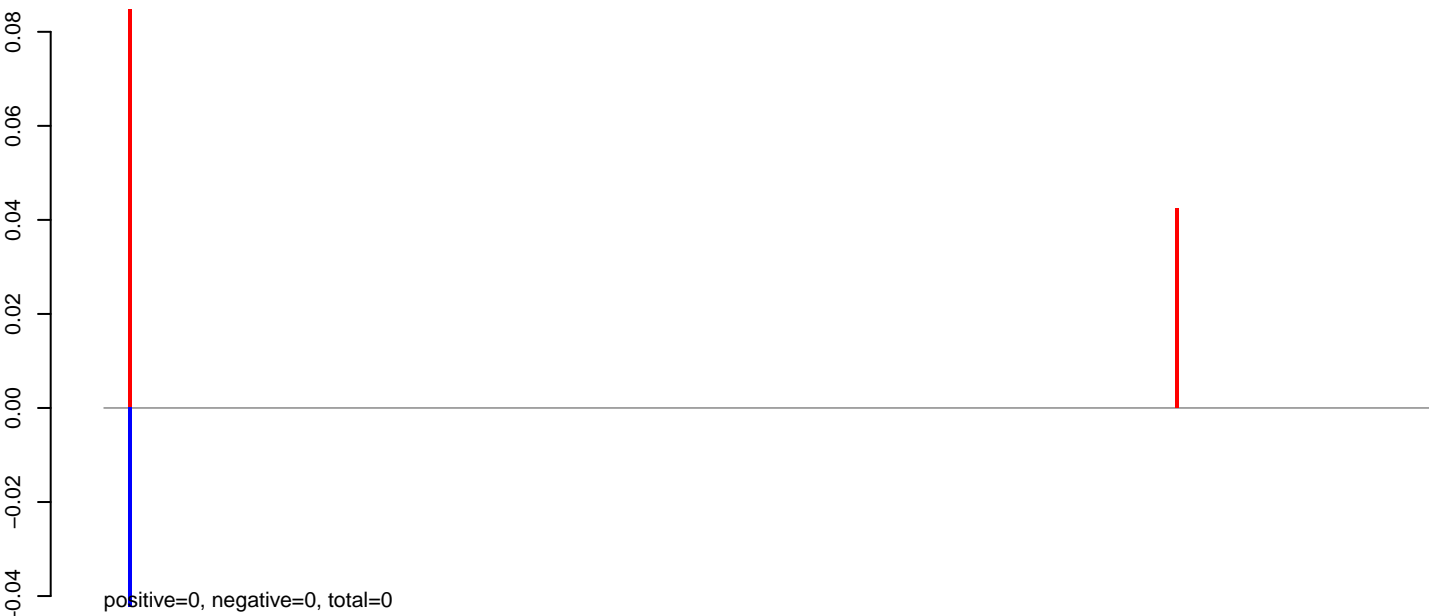
AeAlbo_C636_cells.rep



Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus1-10755

0 2000 4000 6000 8000 10000

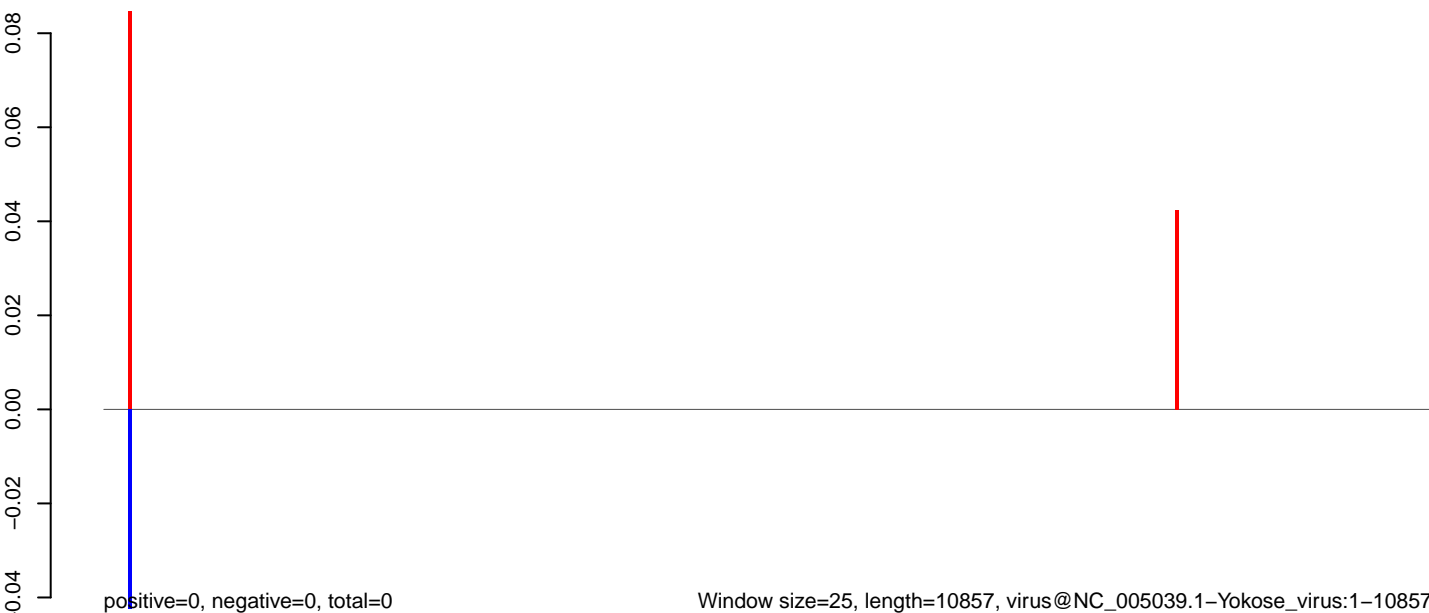
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



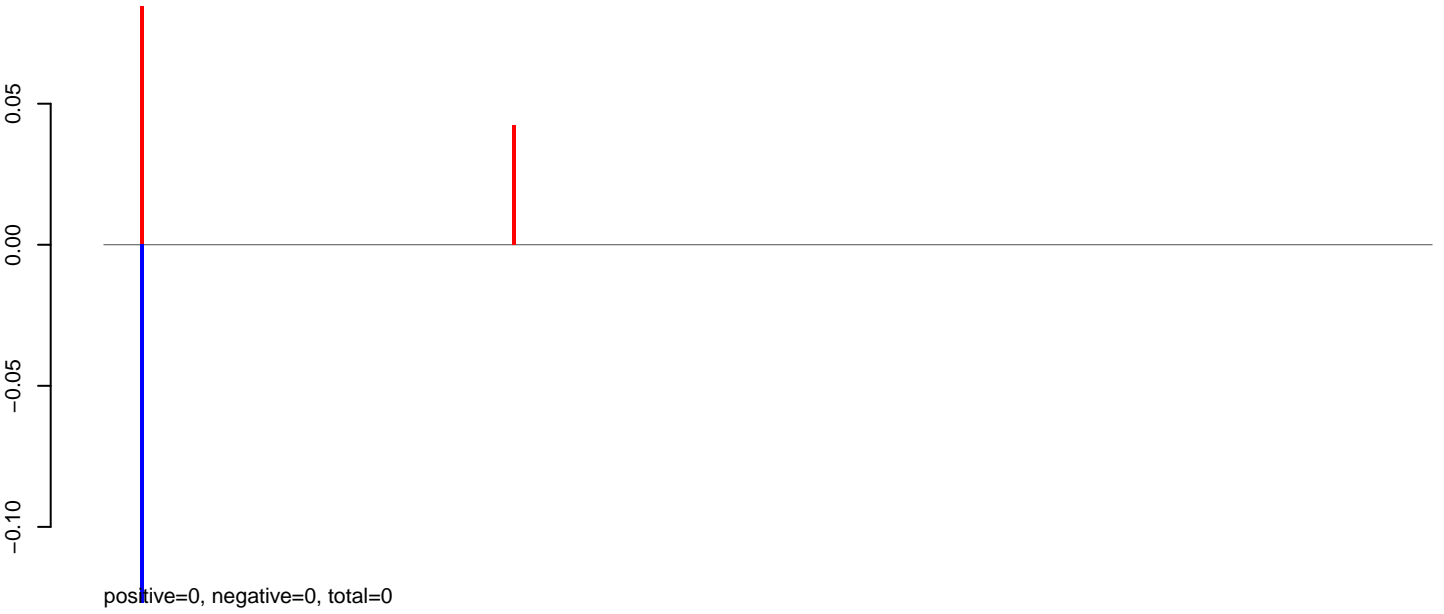
AeAlbo_C636_cells.rep



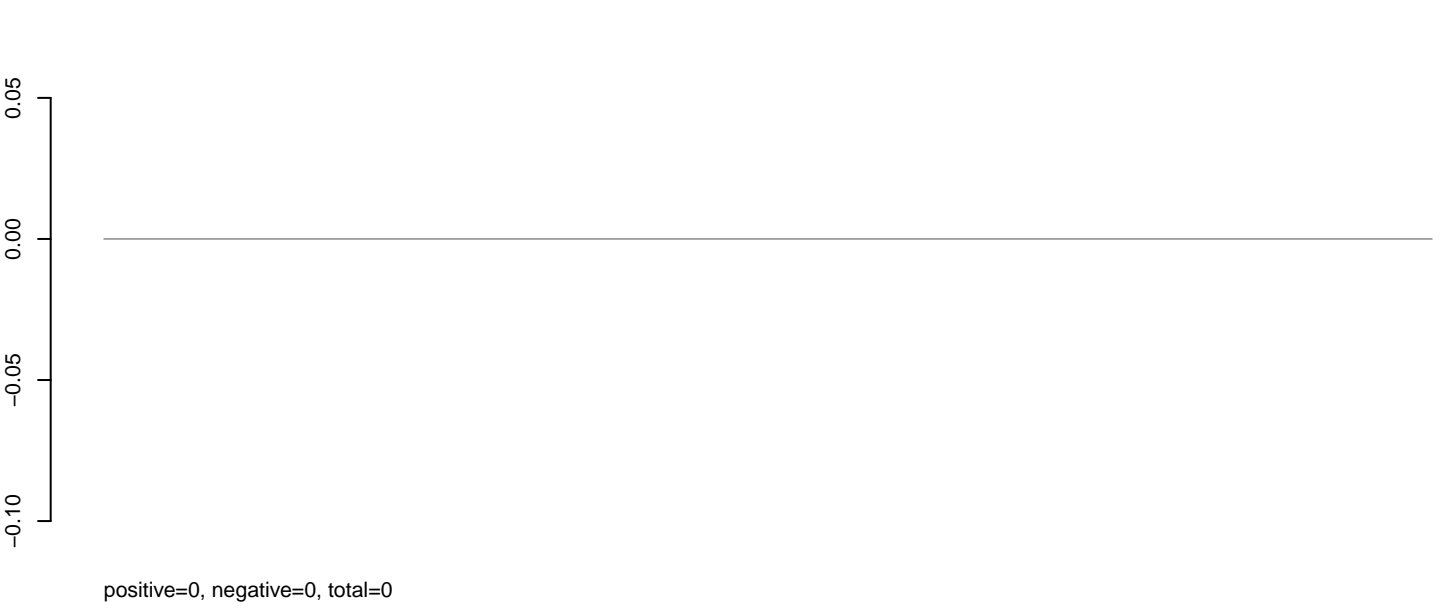
Window size=25, length=10857, virus@NC_005039.1-Yokose_virus:1-10857

0 2000 4000 6000 8000 10000

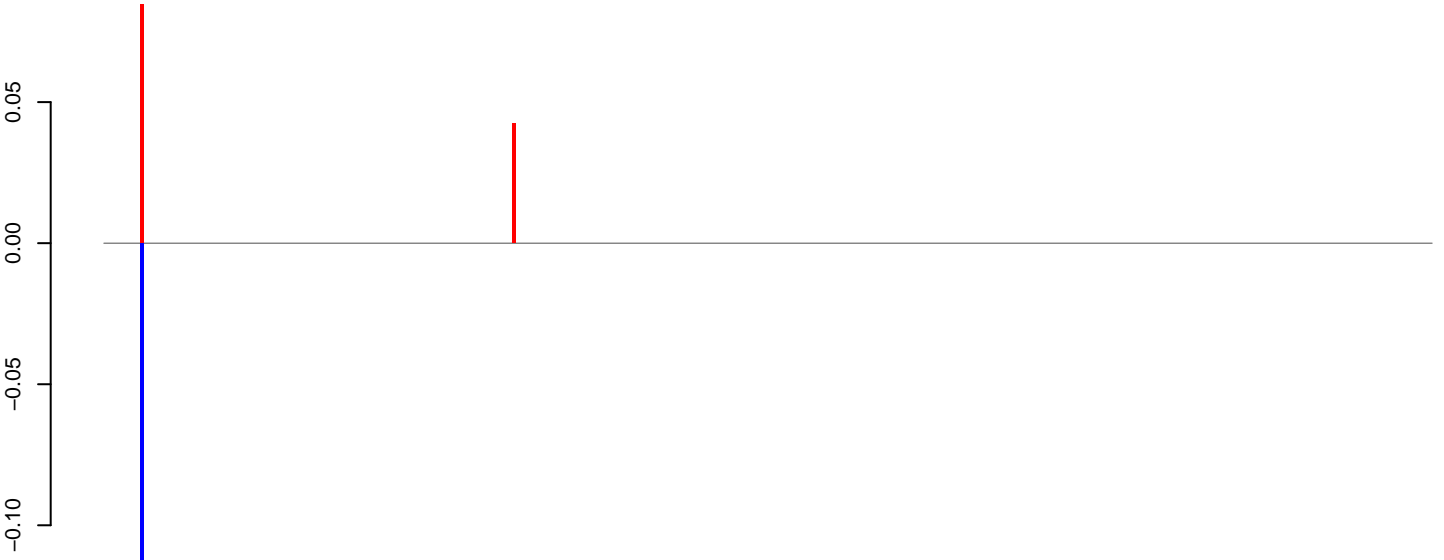
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



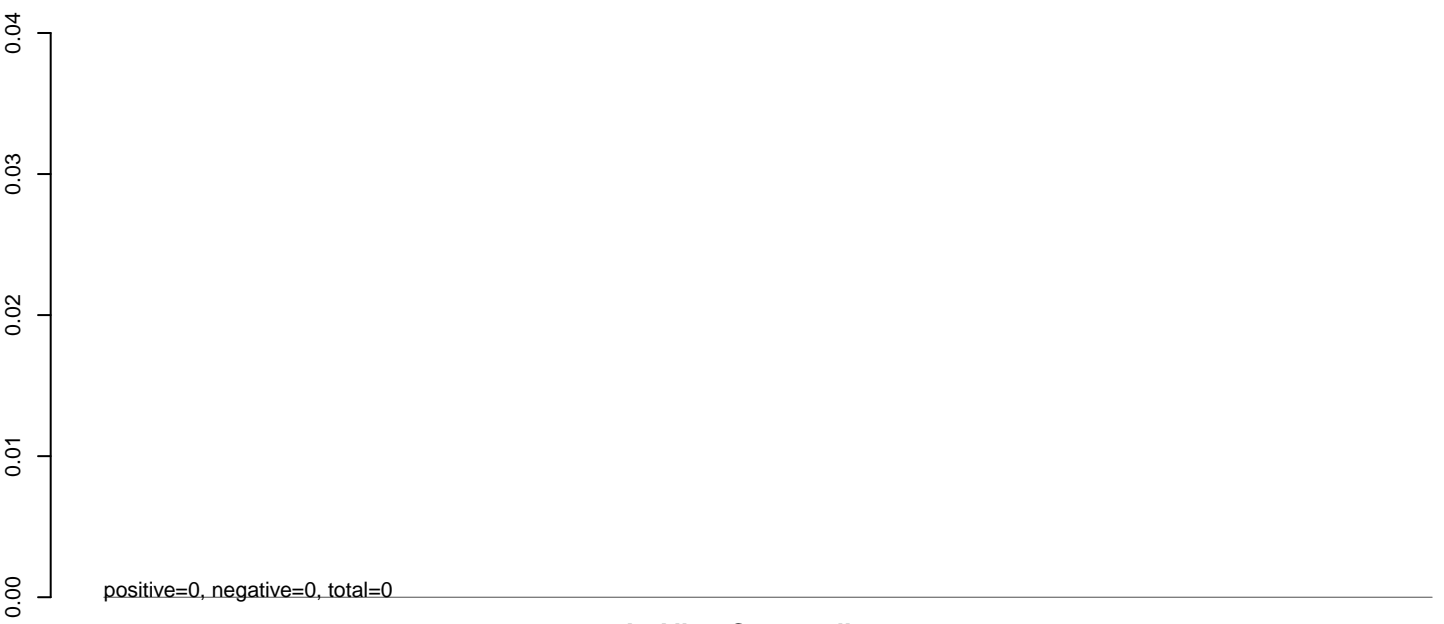
Vindor positive=25, negative=185, total=210 NC_003924.1-Cricket_paralysis_virus_nonstructural_polyprotein_and_structural_polyprotein_genes:1-9185

0 2000 4000 6000 8000

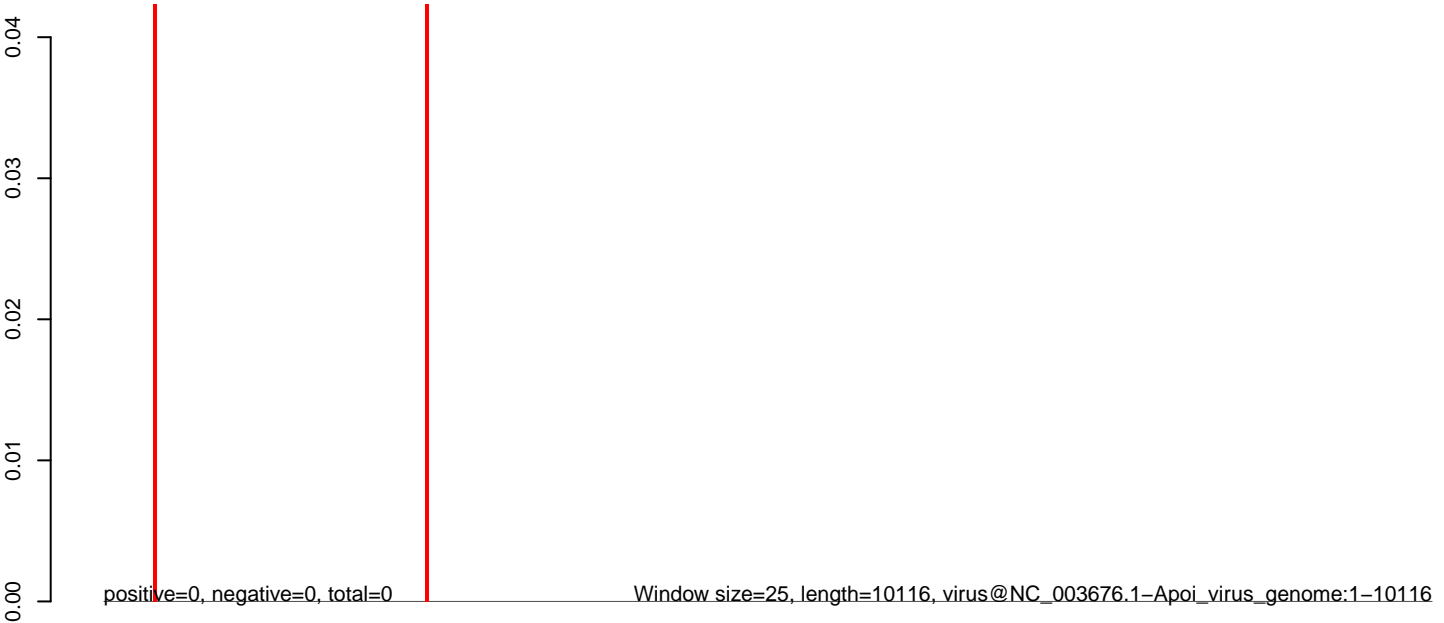
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

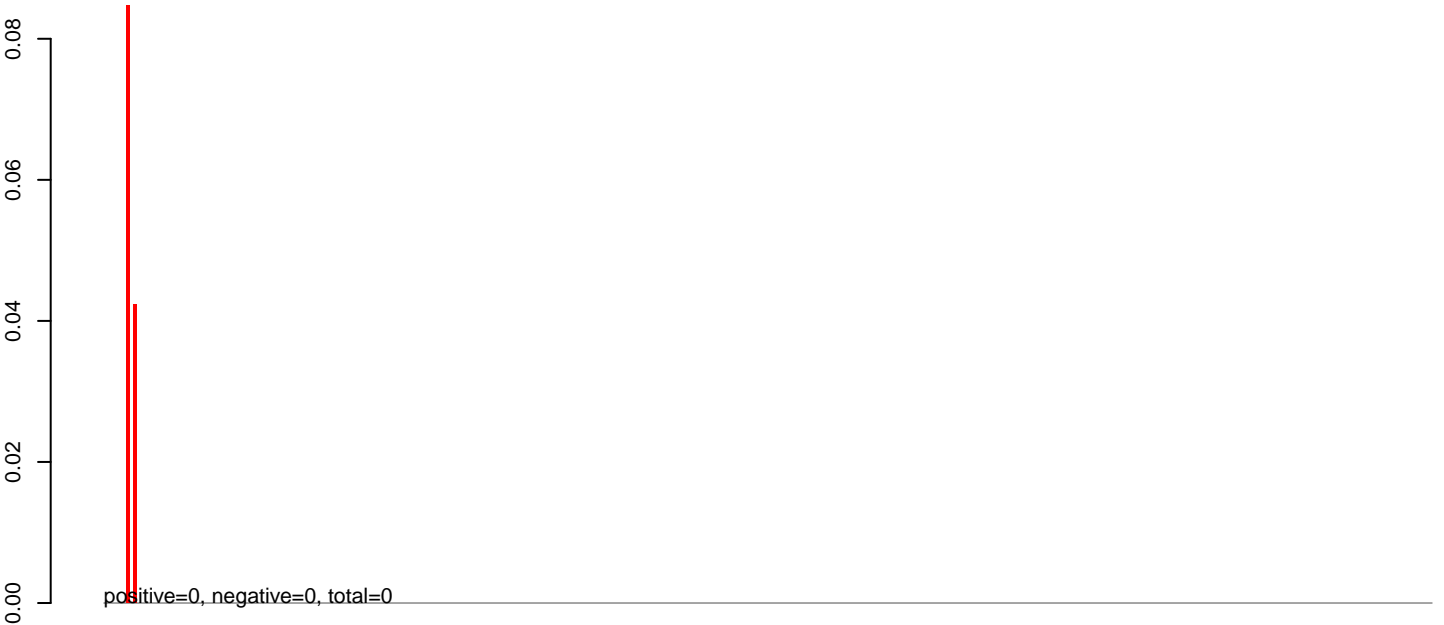


AeAlbo_C636_cells.rep

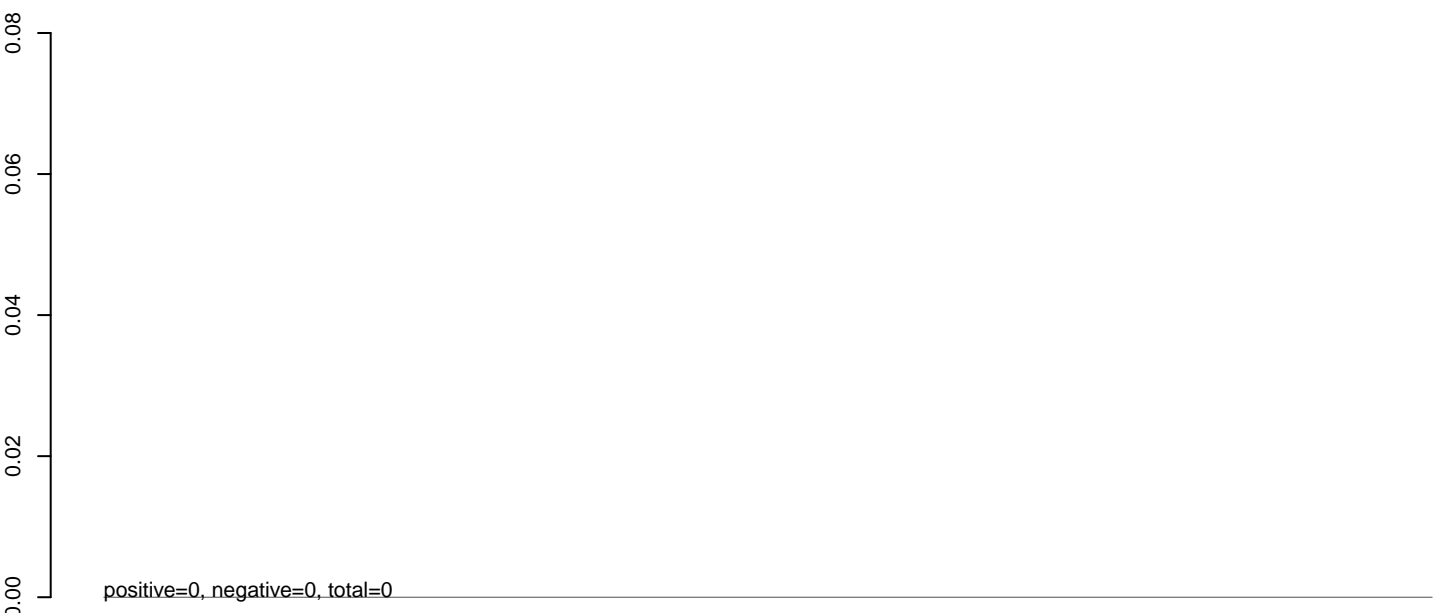


0 2000 4000 6000 8000 10000

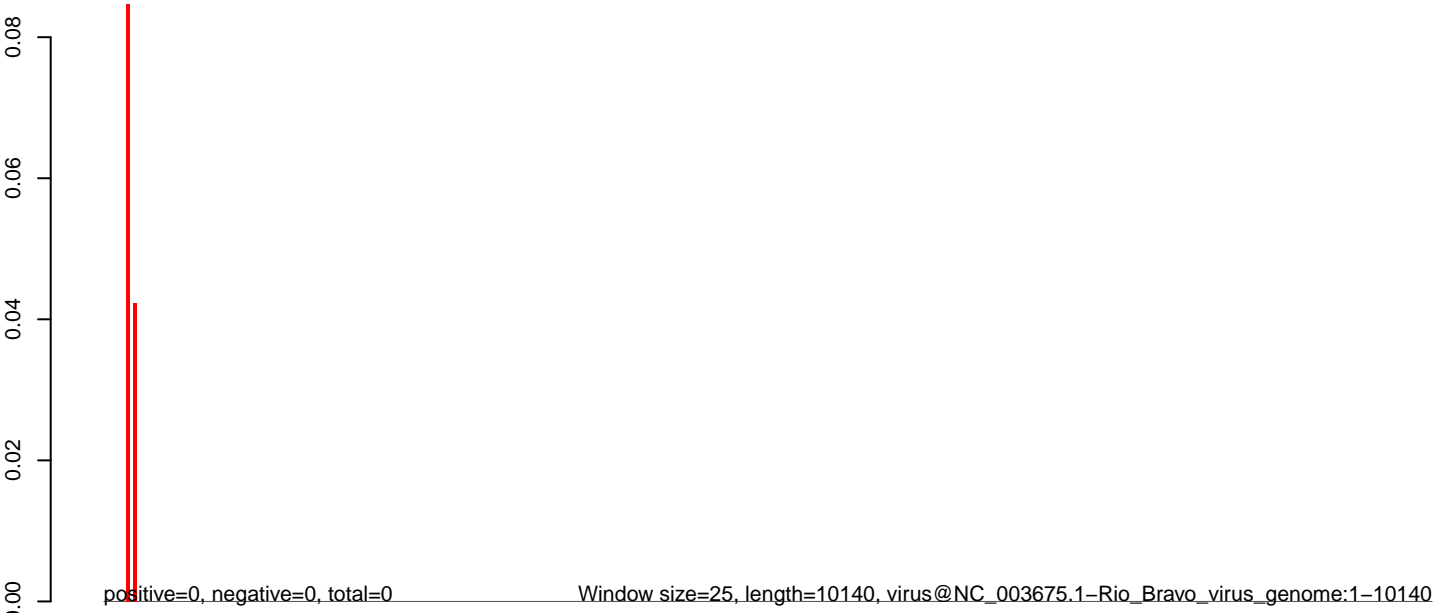
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



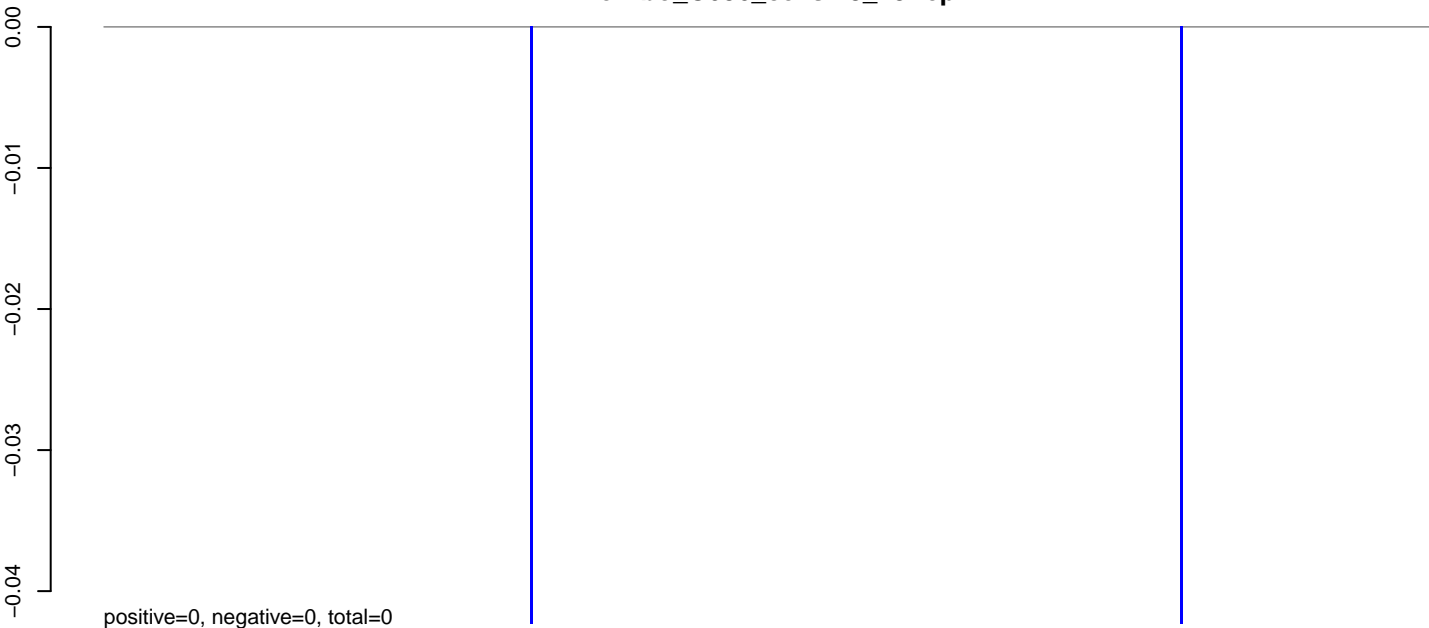
AeAlbo_C636_cells.rep



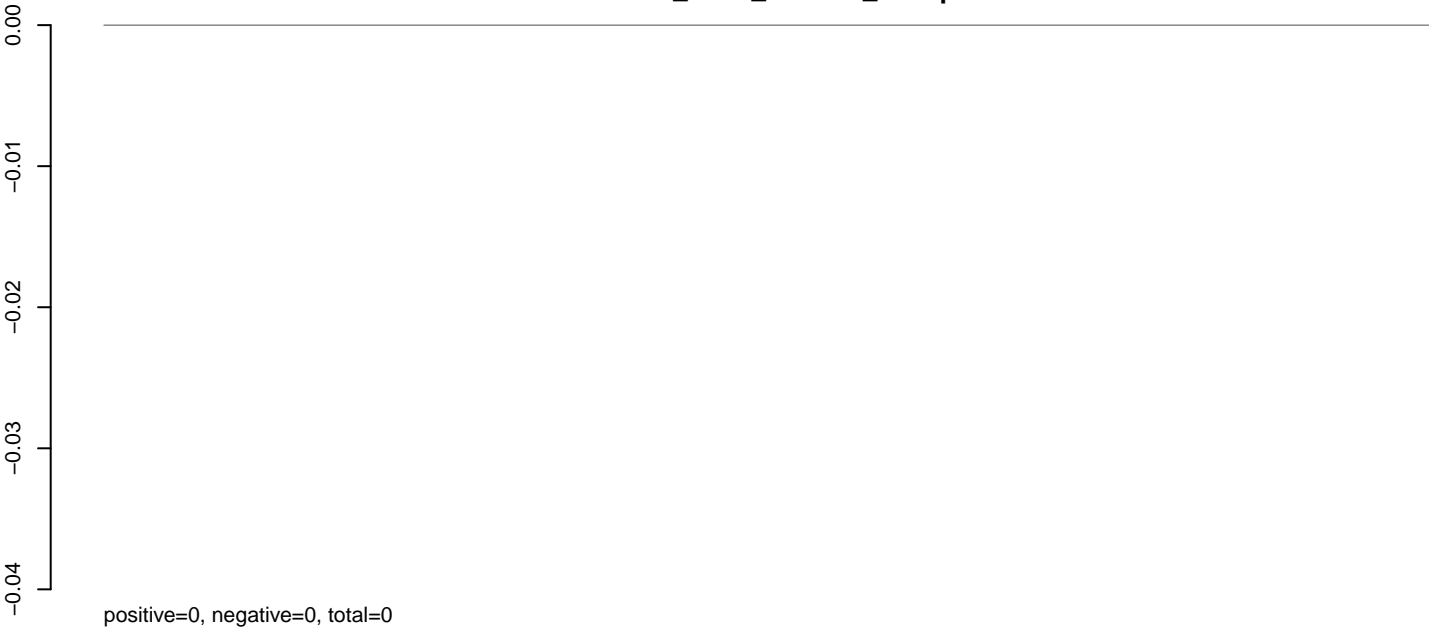
Window size=25, length=10140, virus@NC_003675.1-Rio_Bravo_virus_genome:1-10140

0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

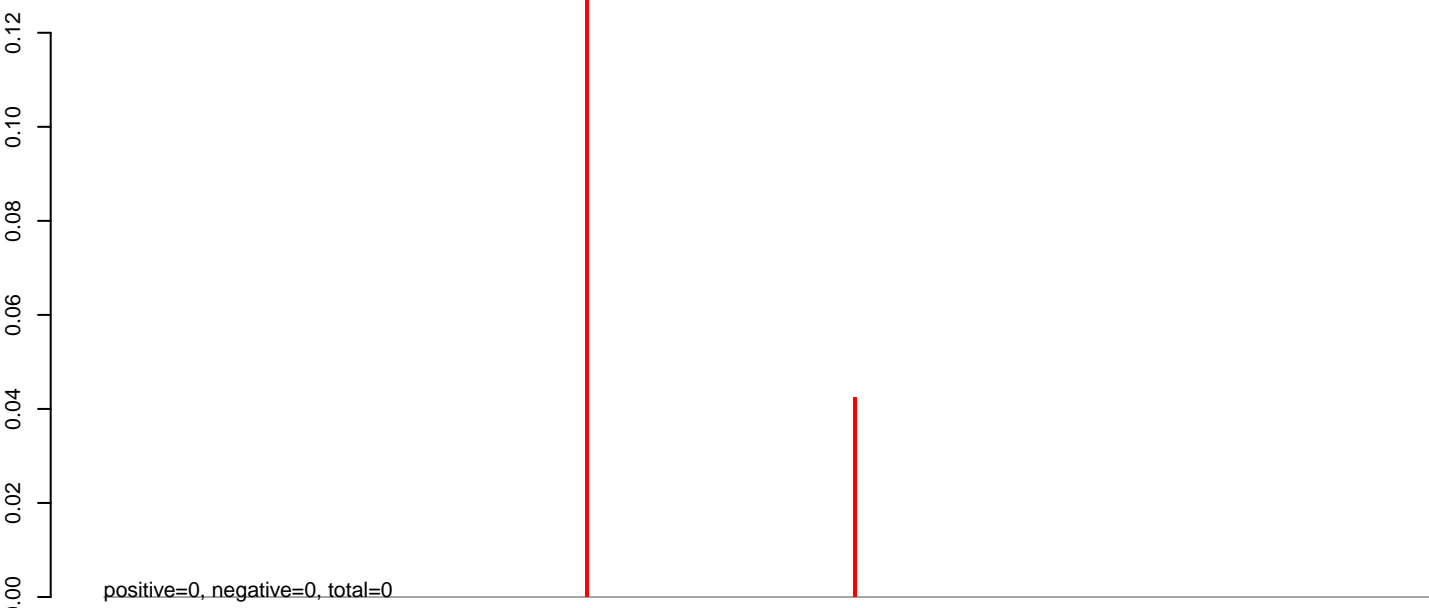


AeAlbo_C636_cells.rep

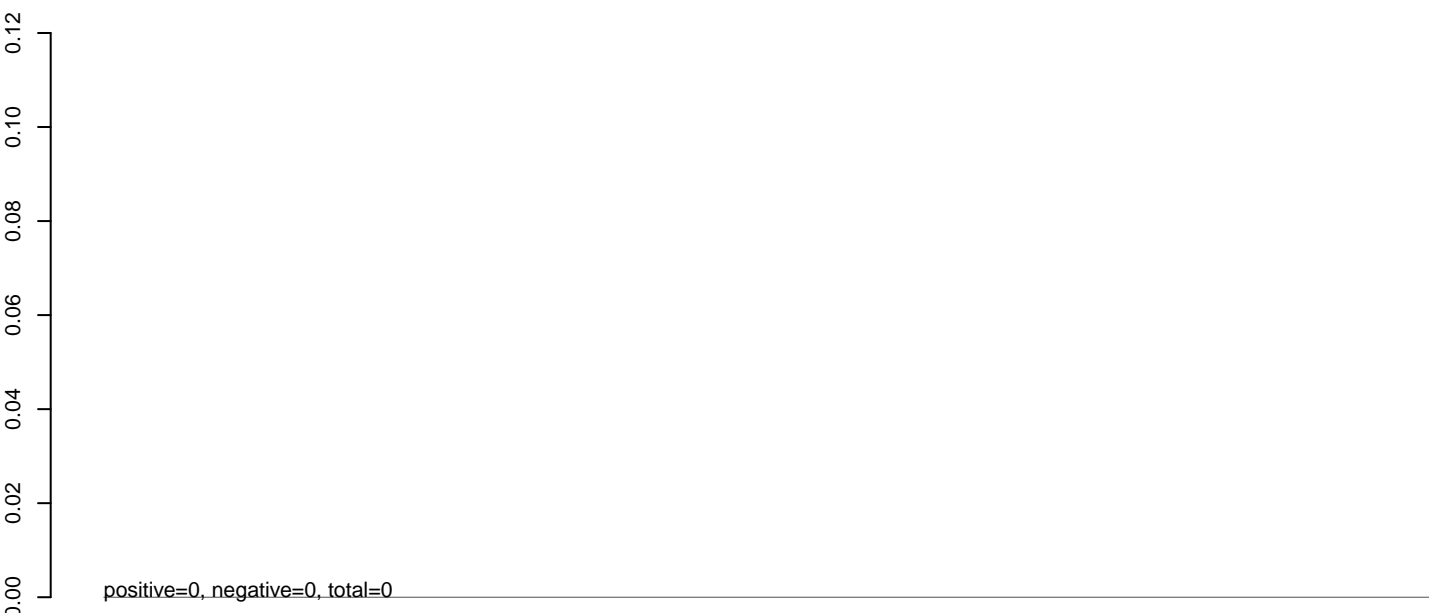


0 2000 4000 6000 8000 10000

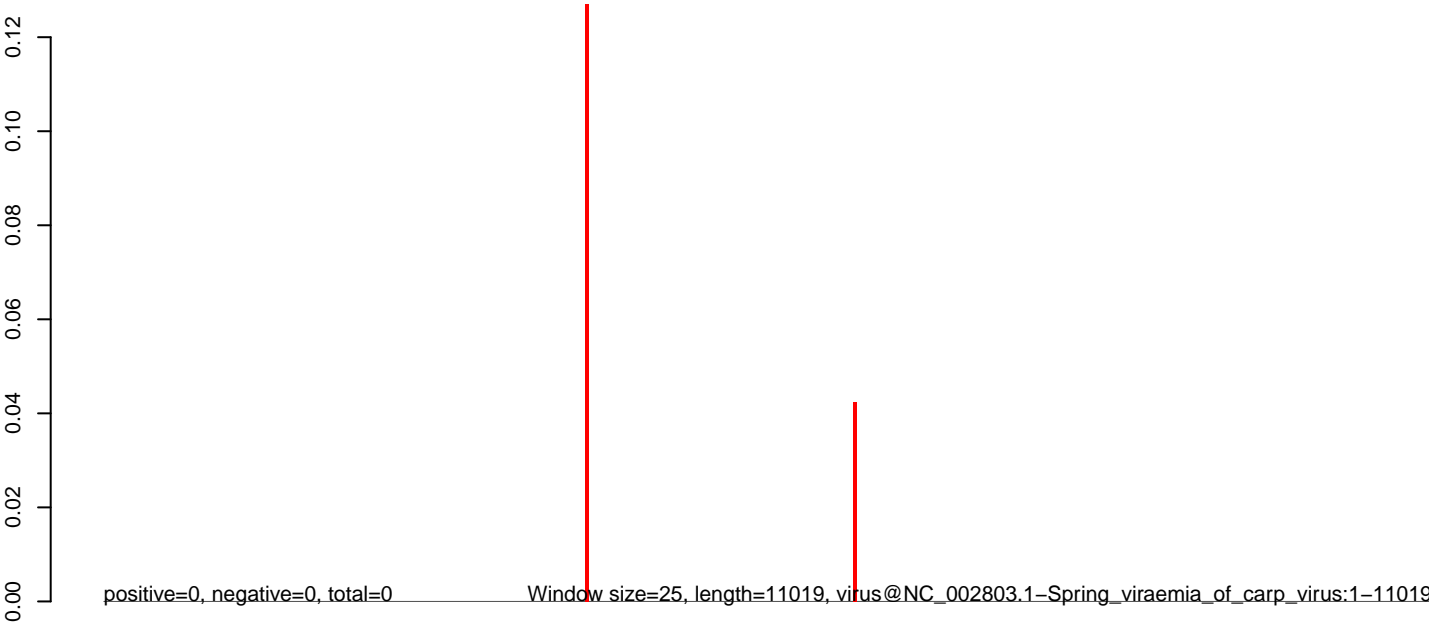
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

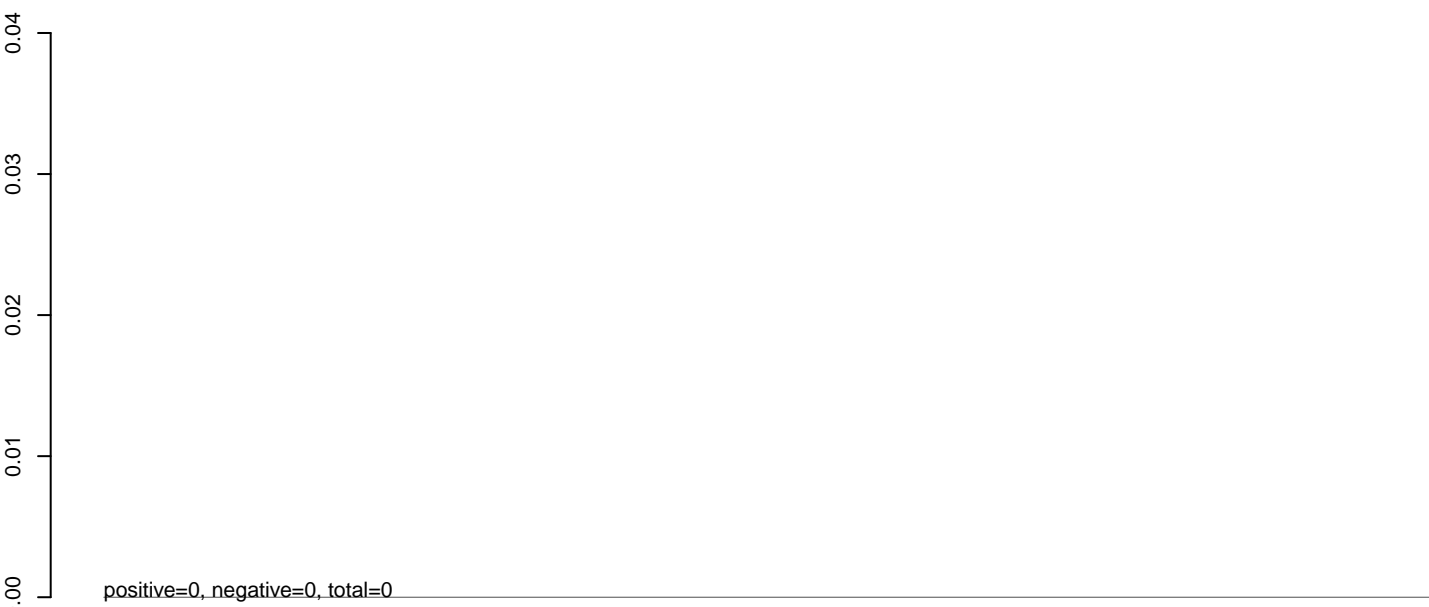


0 2000 4000 6000 8000 10000

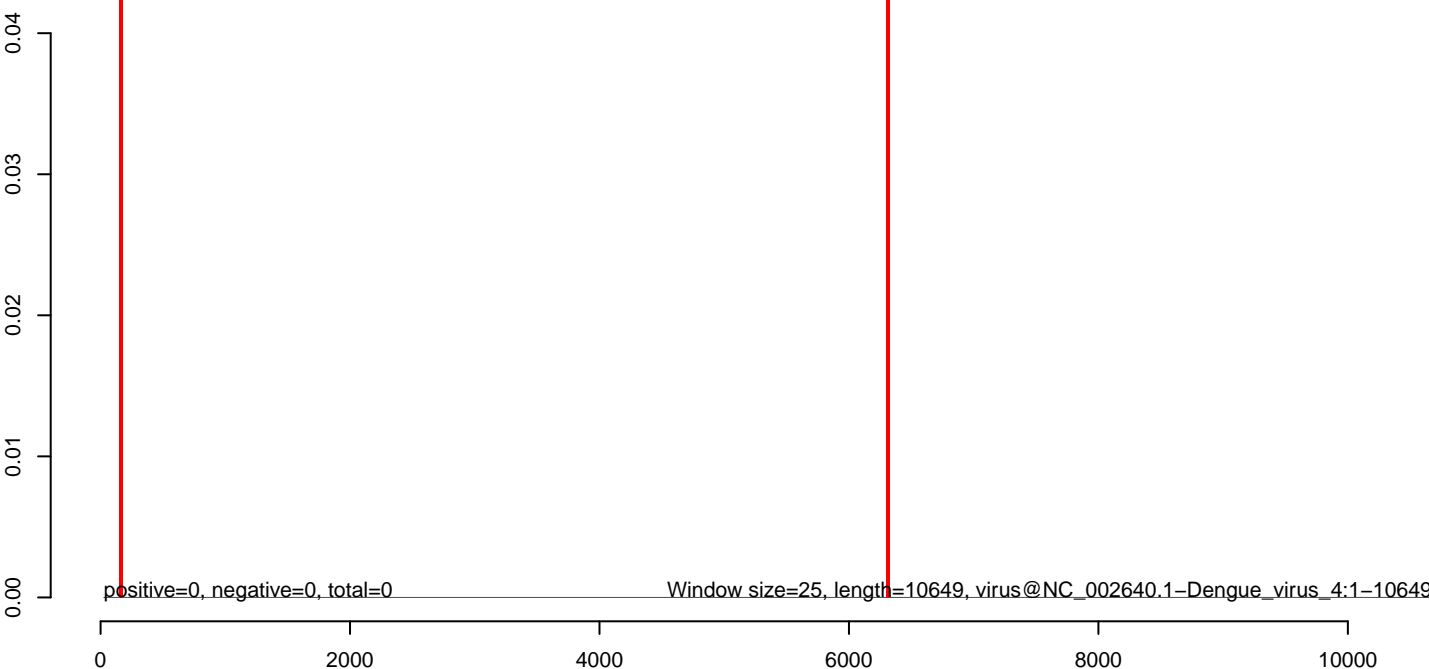
AeAlbo_C636_cells.18_23.rep



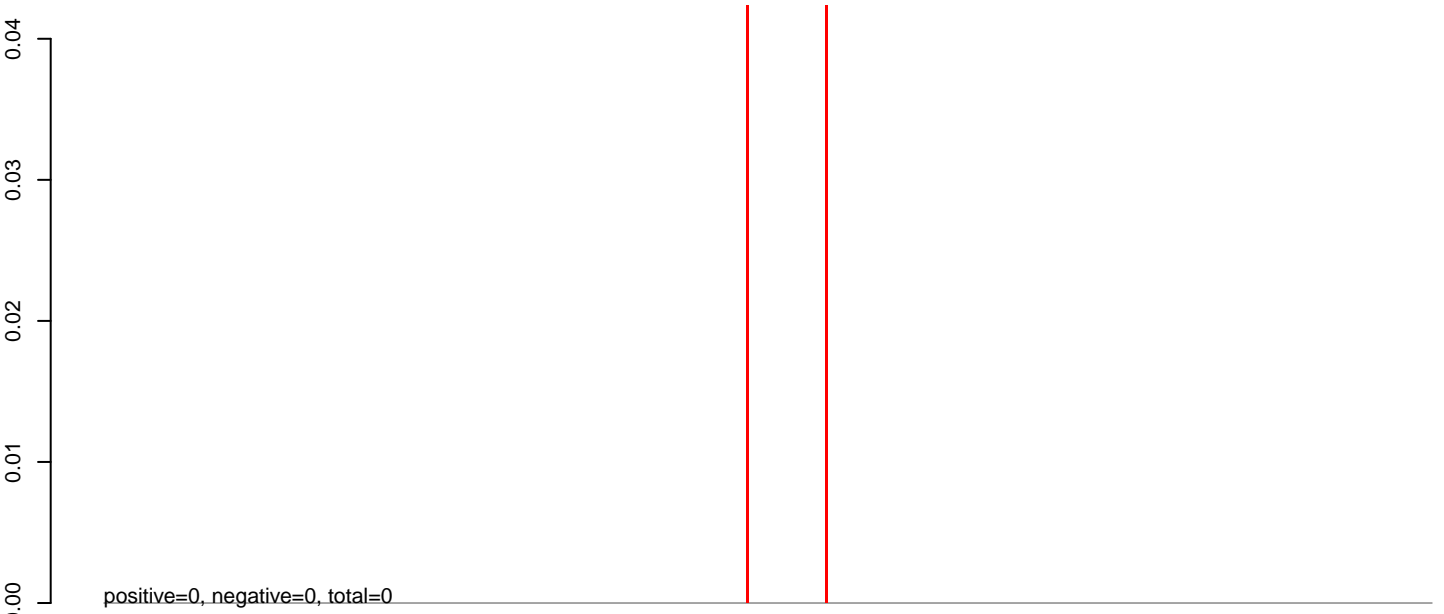
AeAlbo_C636_cells.24_35.rep



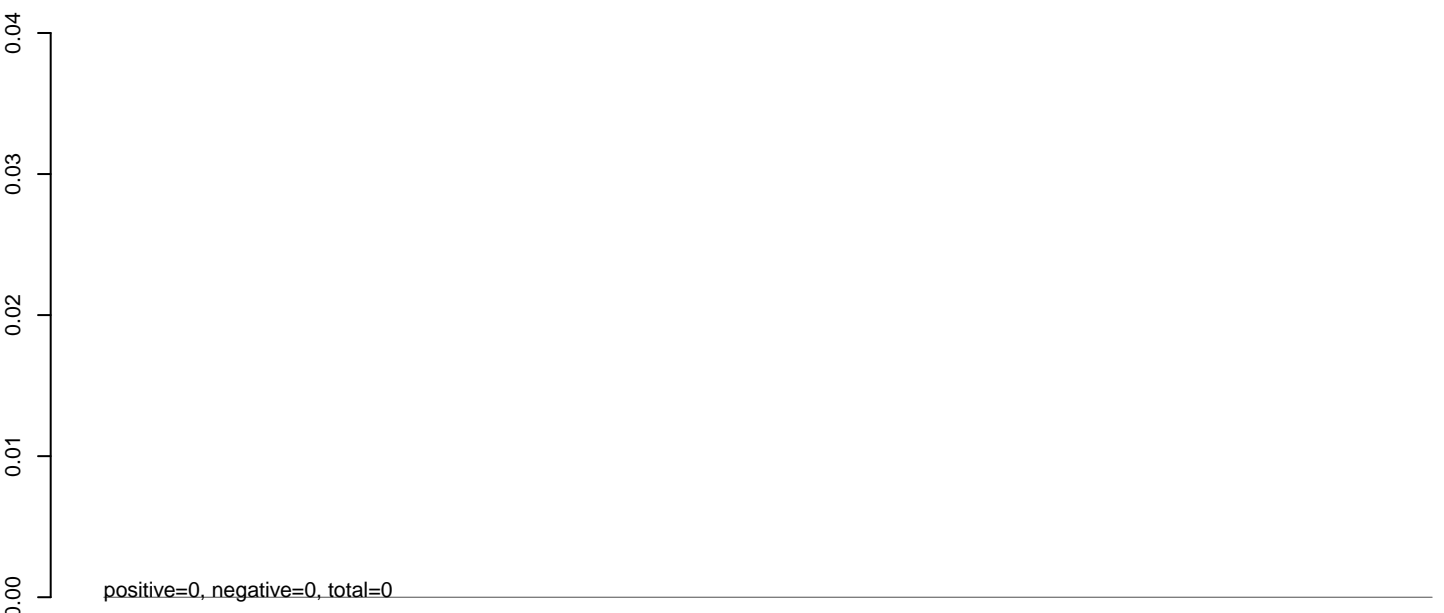
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

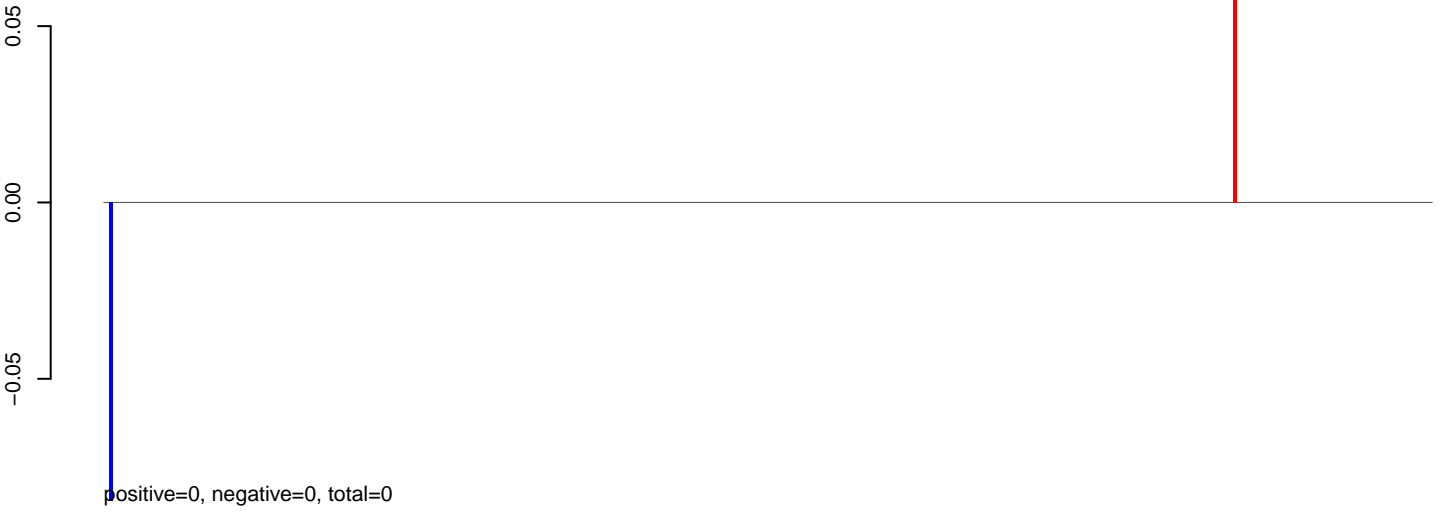


AeAlbo_C636_cells.rep

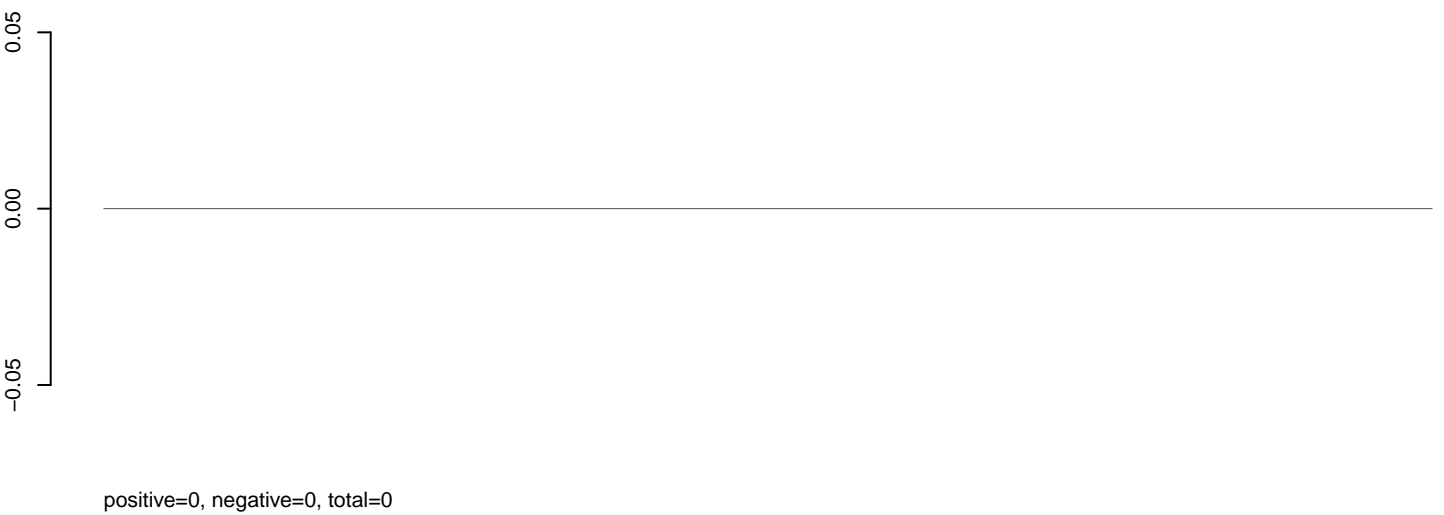


0 2000 4000 6000 8000 10000 12000
positive=0, negative=0, total=0
window size=25, length=12294, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

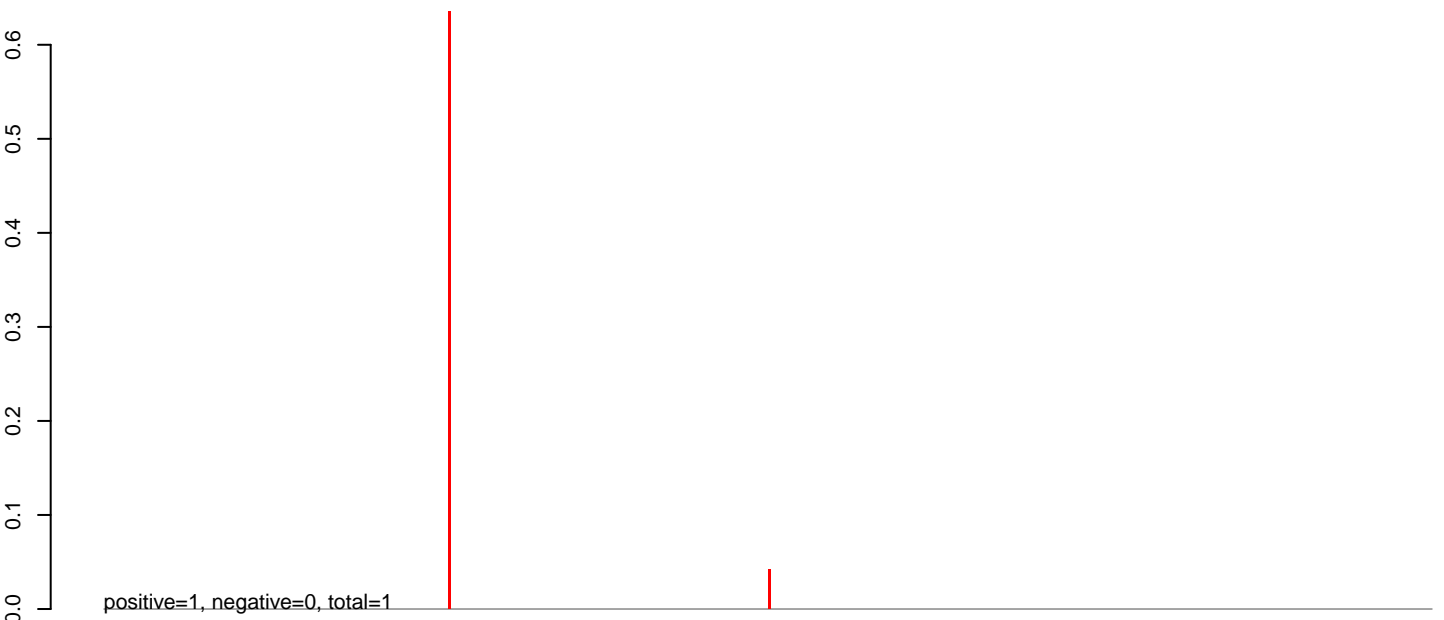


AeAlbo_C636_cells.rep

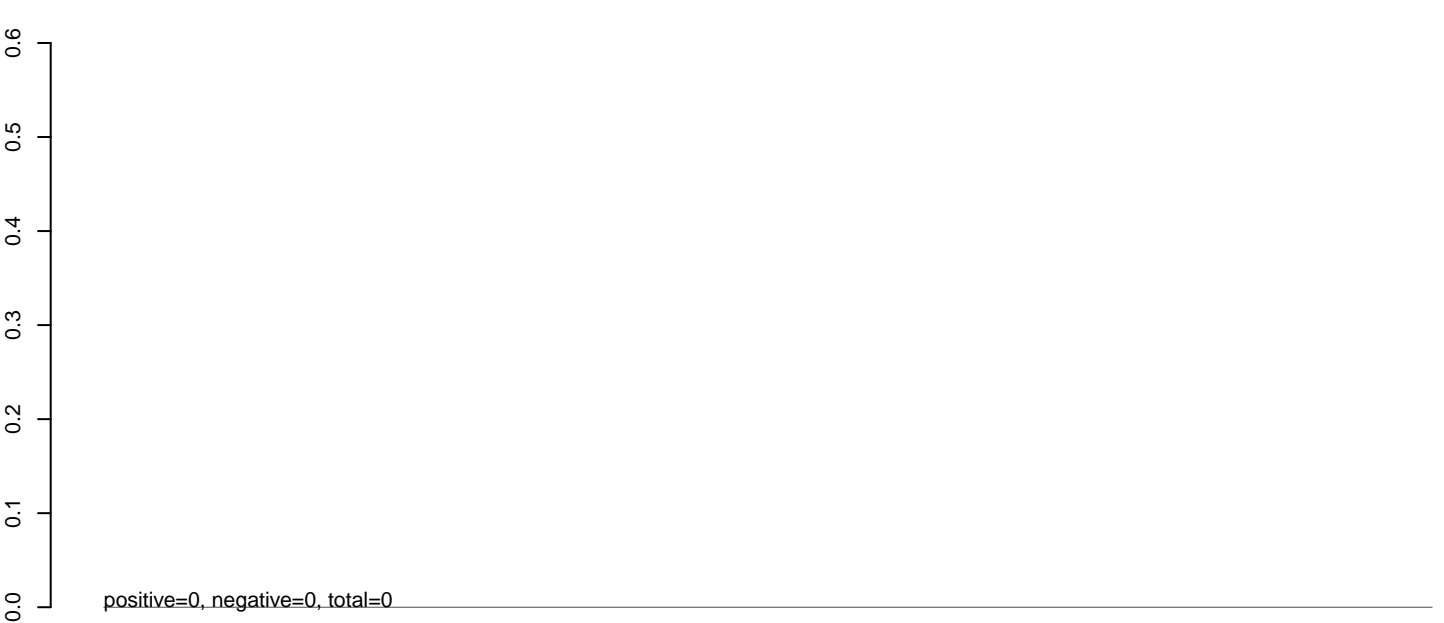


0 2000 4000 6000 8000 10000

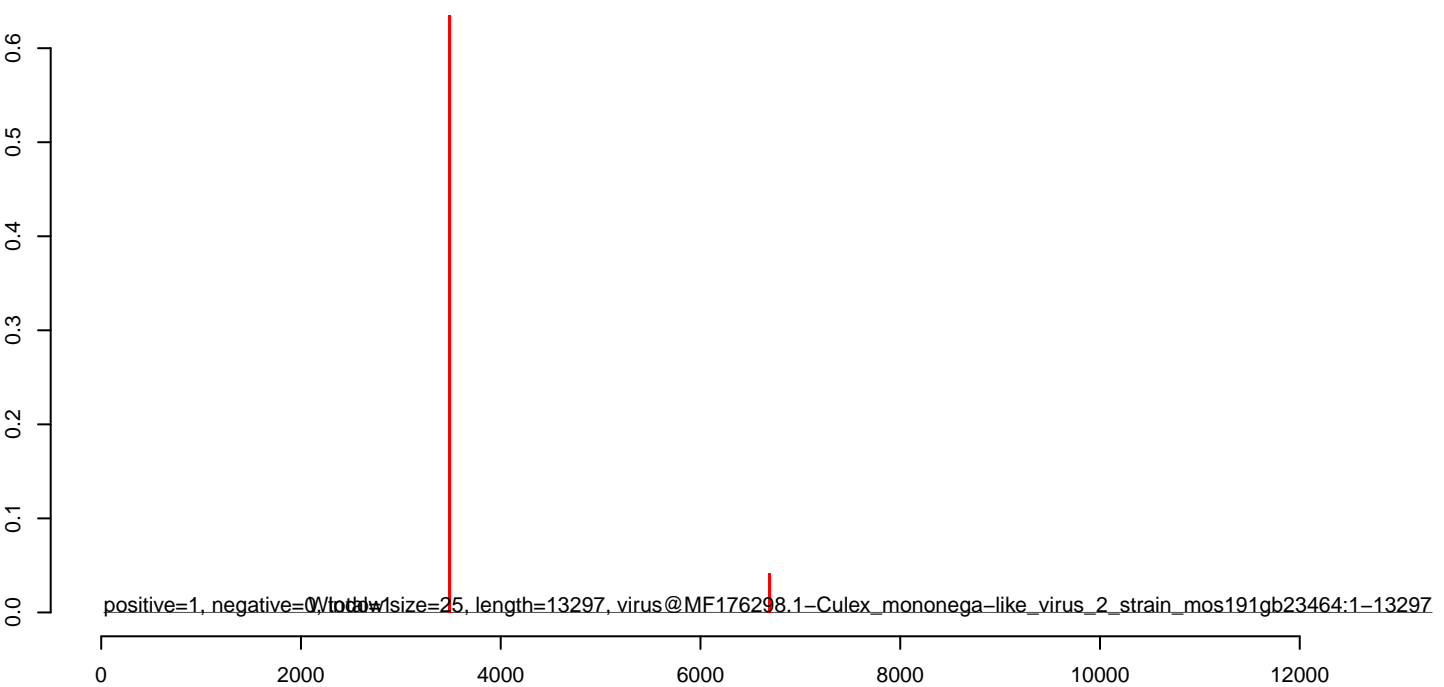
AeAlbo_C636_cells.18_23.rep



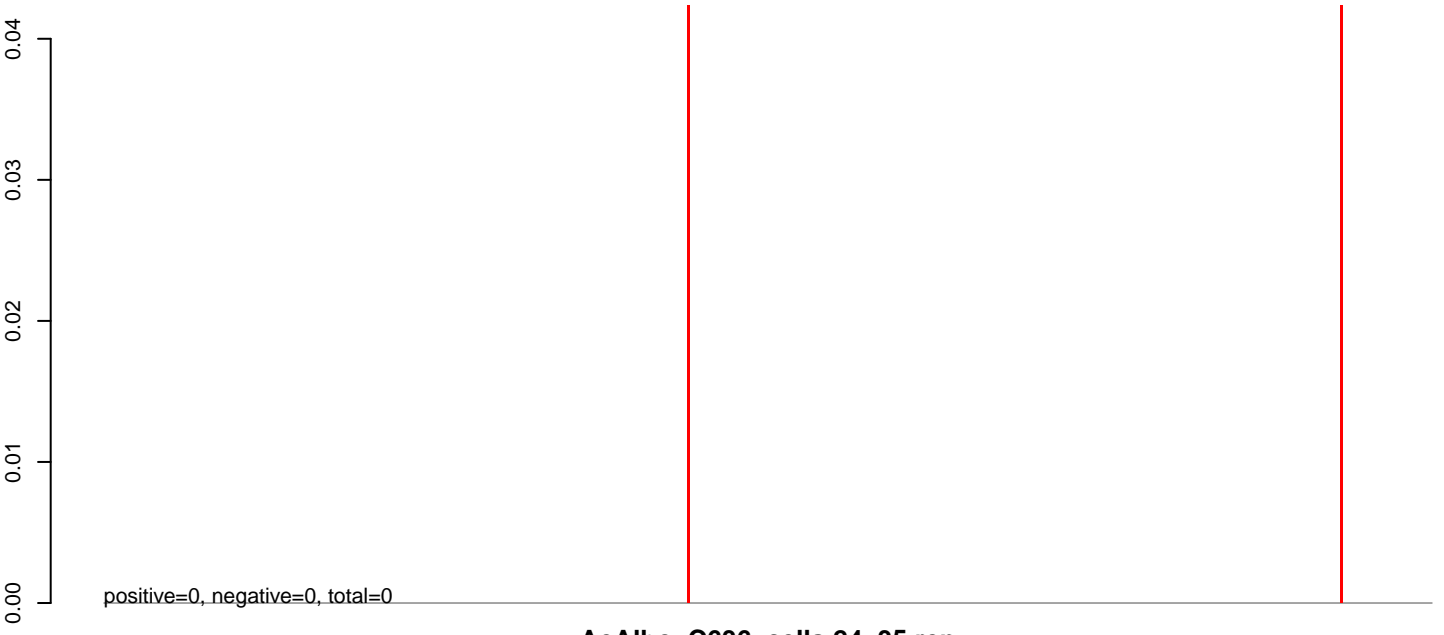
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



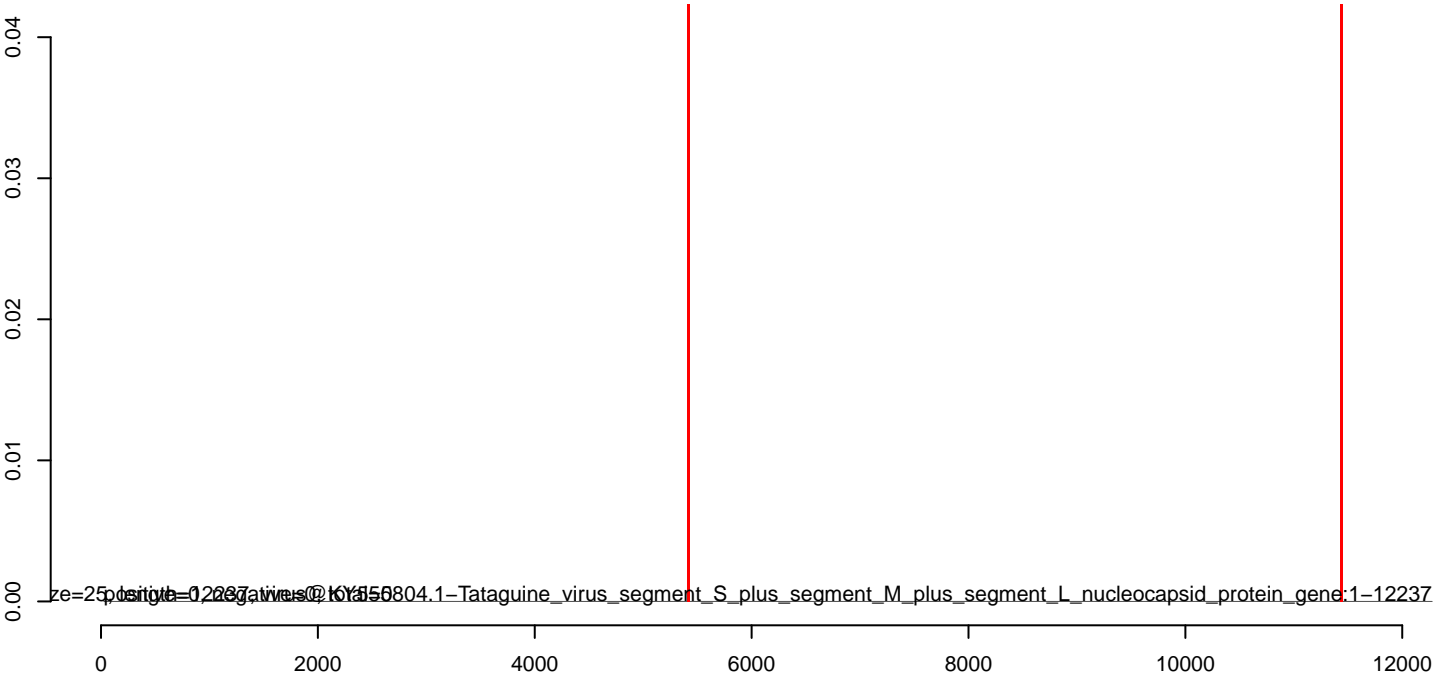
AeAlbo_C636_cells.18_23.rep



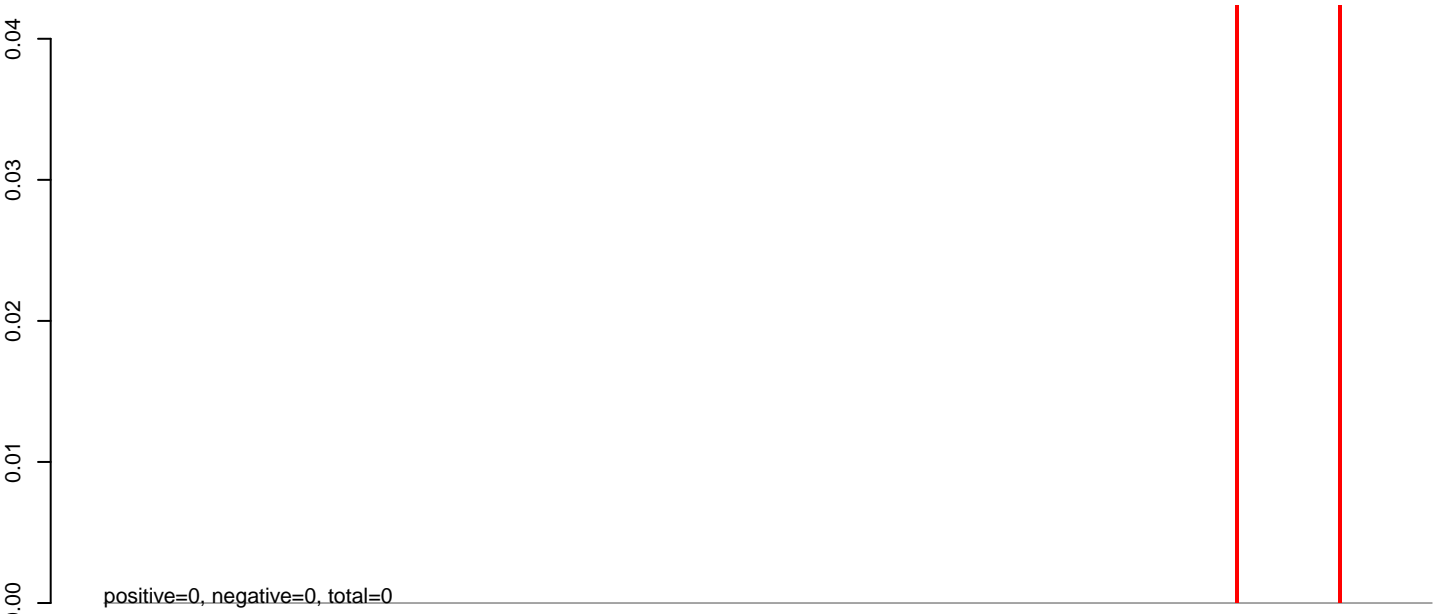
AeAlbo_C636_cells.24_35.rep



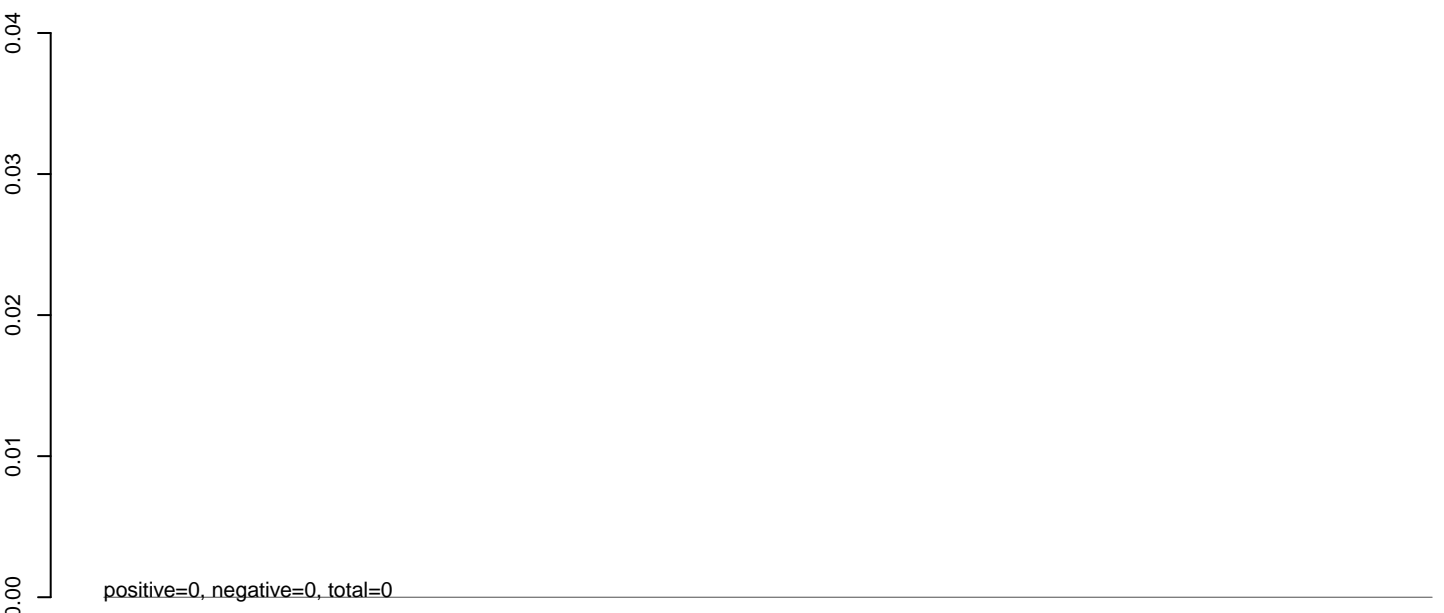
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

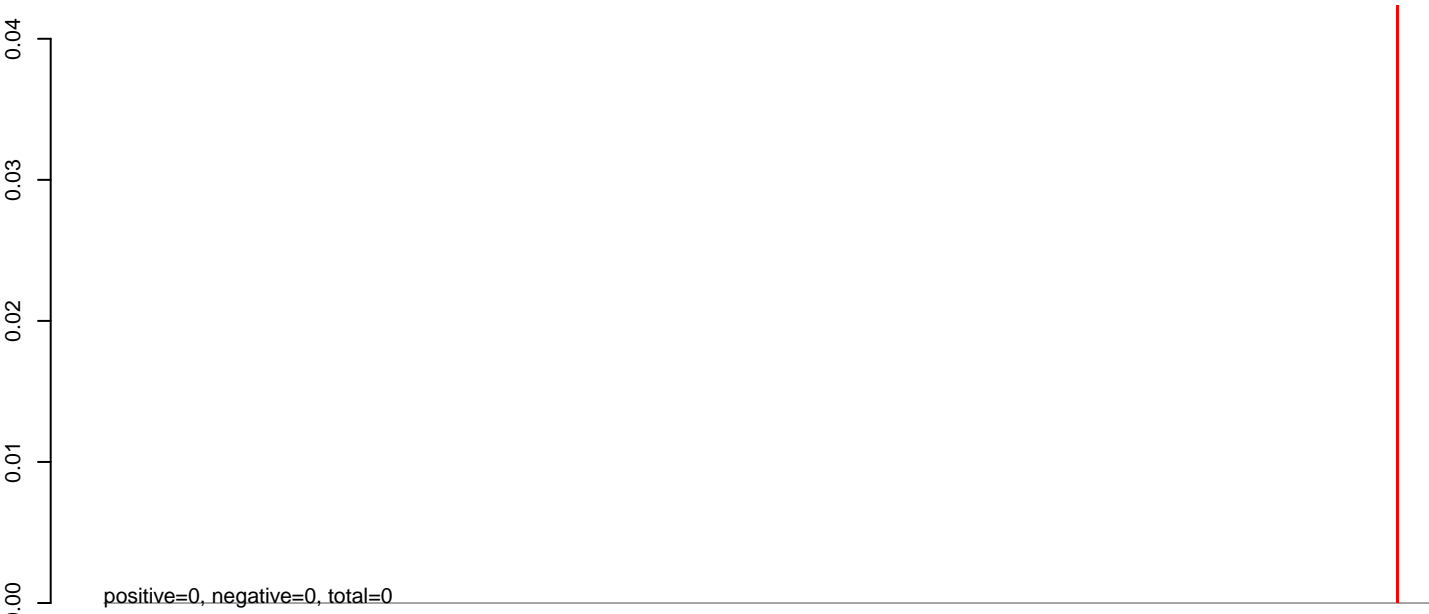


AeAlbo_C636_cells.rep

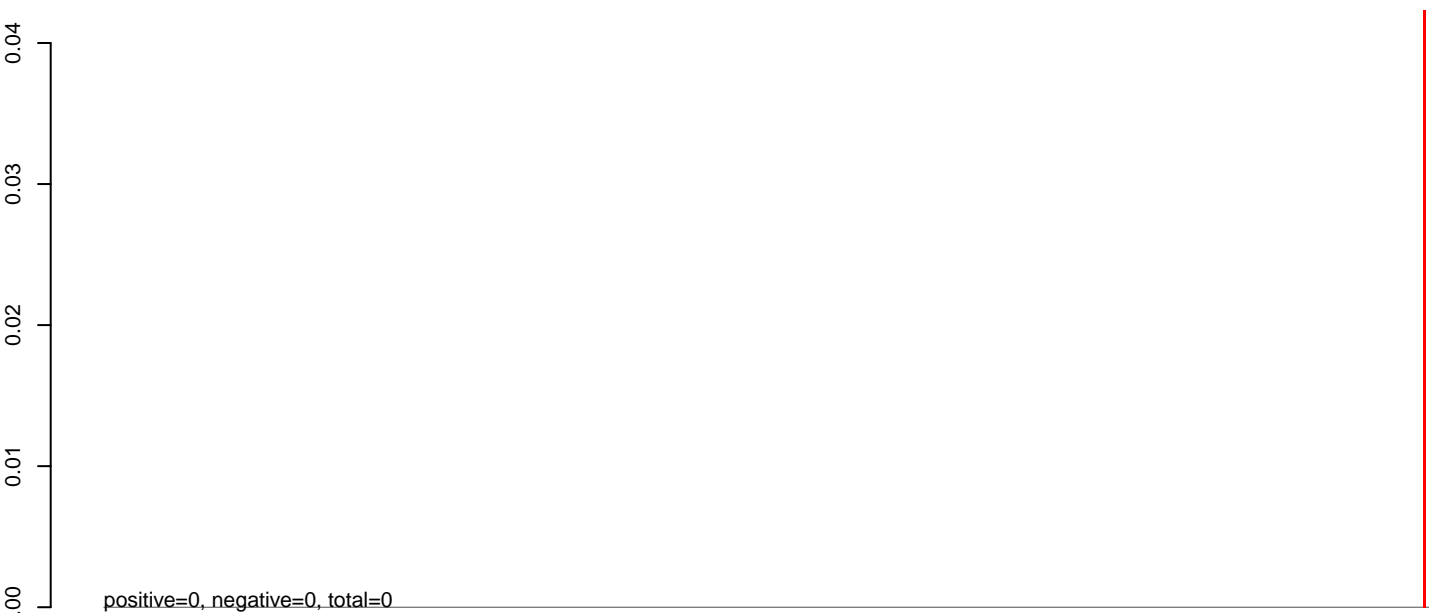


0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



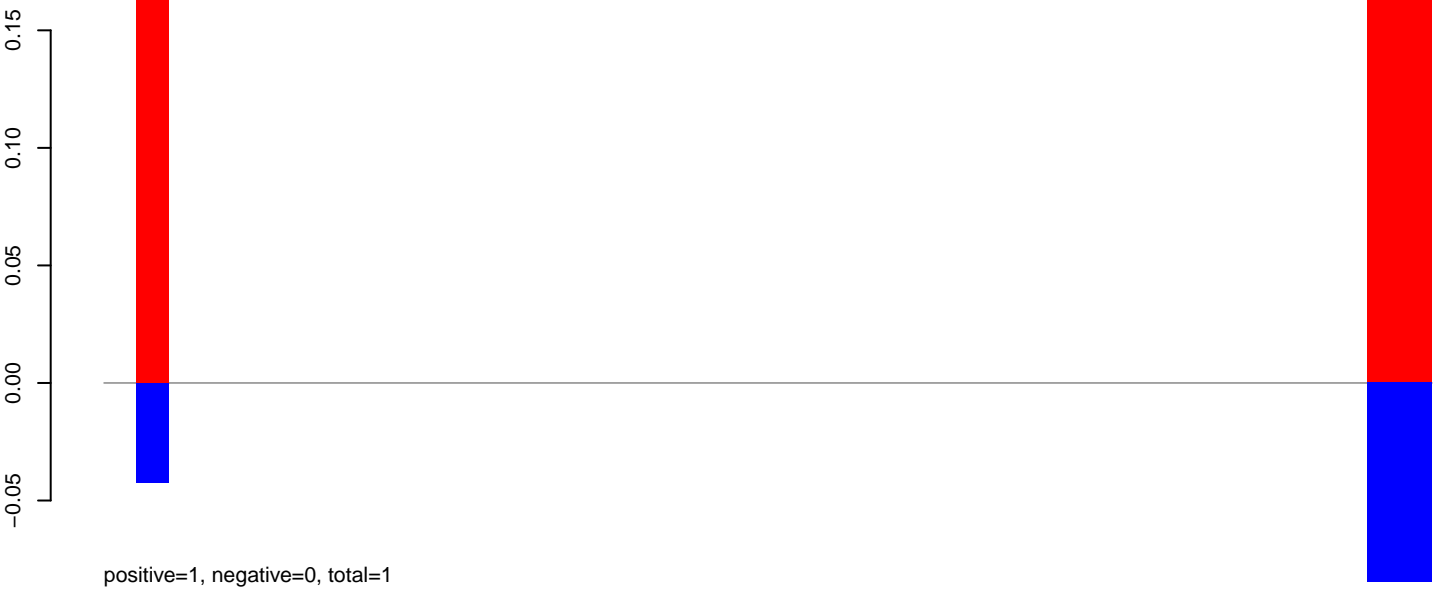
AeAlbo_C636_cells.24_35.rep



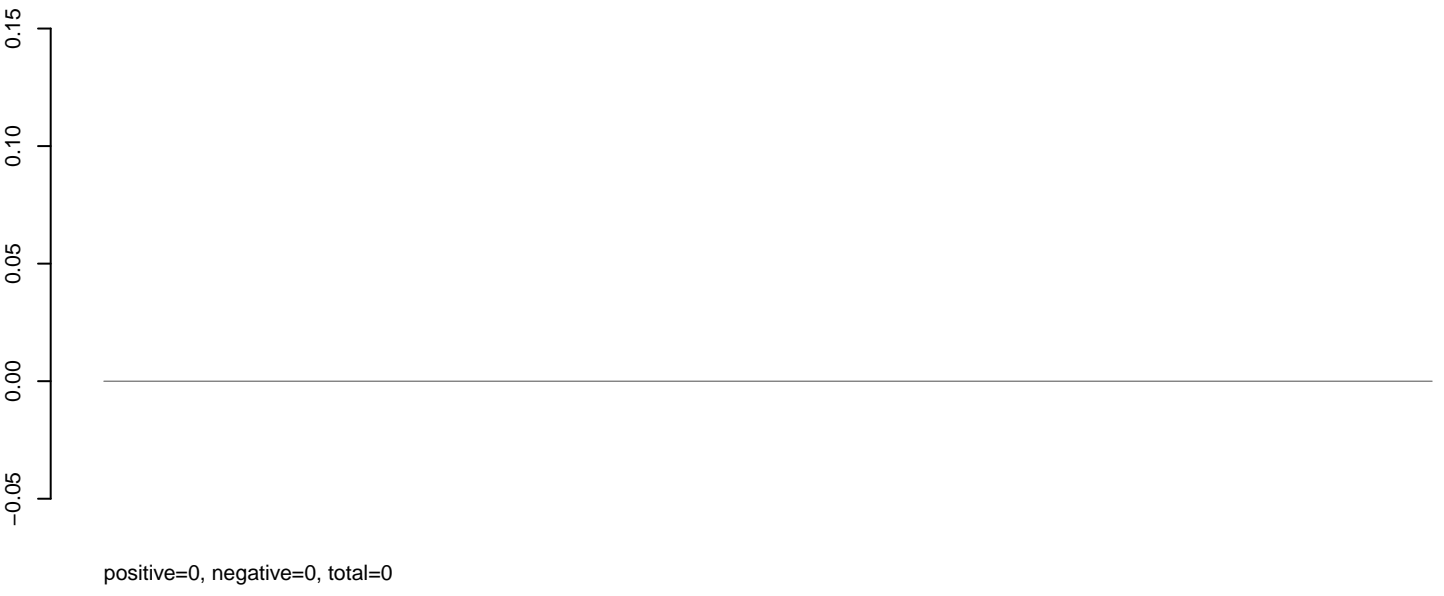
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



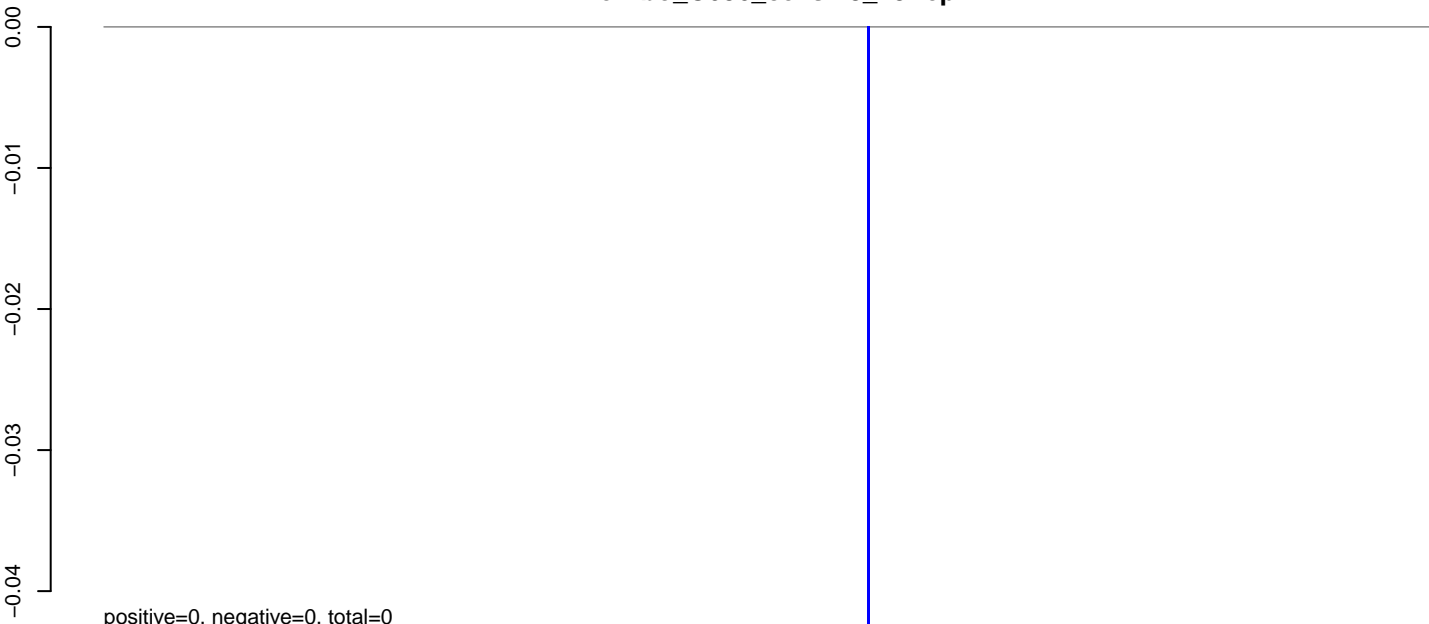
AeAlbo_C636_cells.24_35.rep



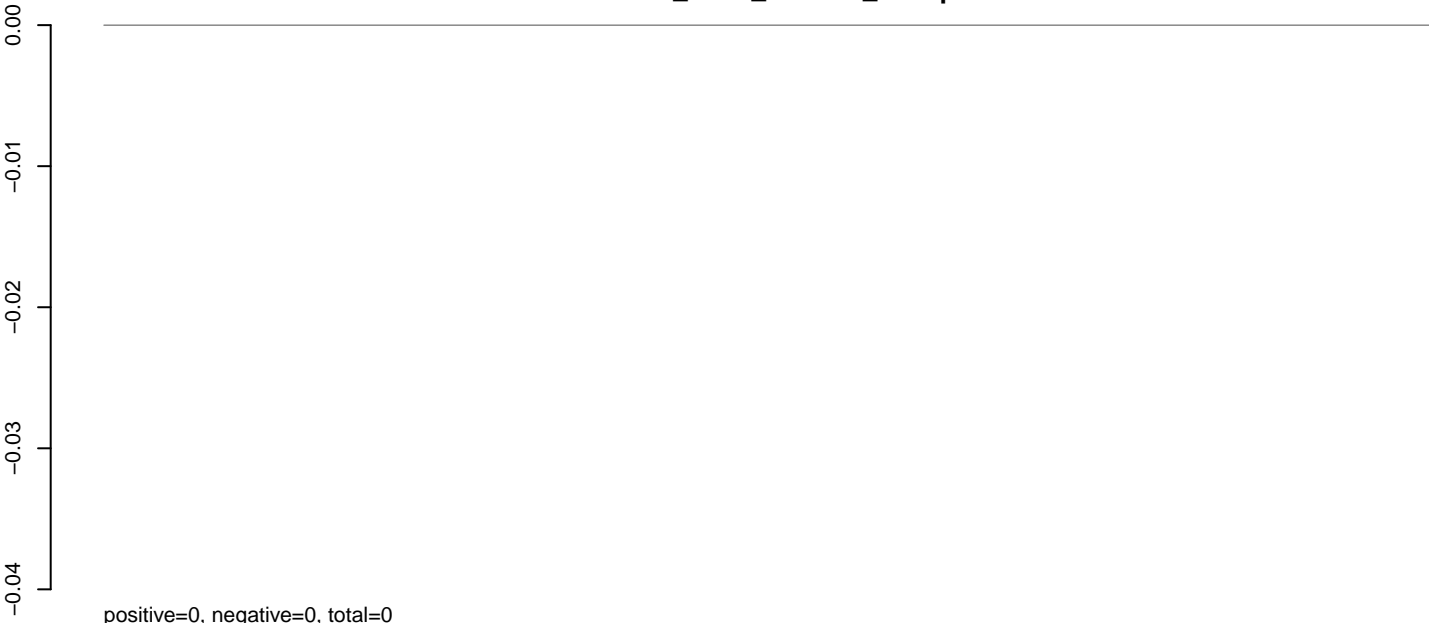
AeAlbo_C636_cells.rep



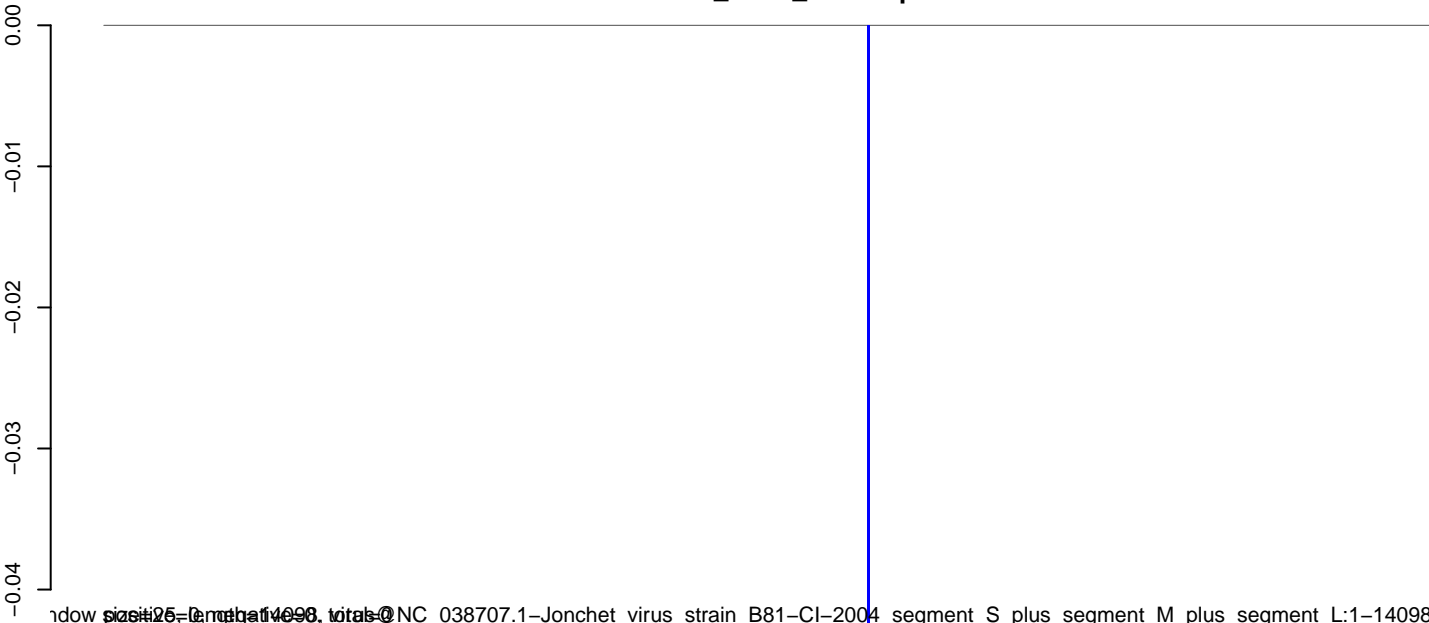
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



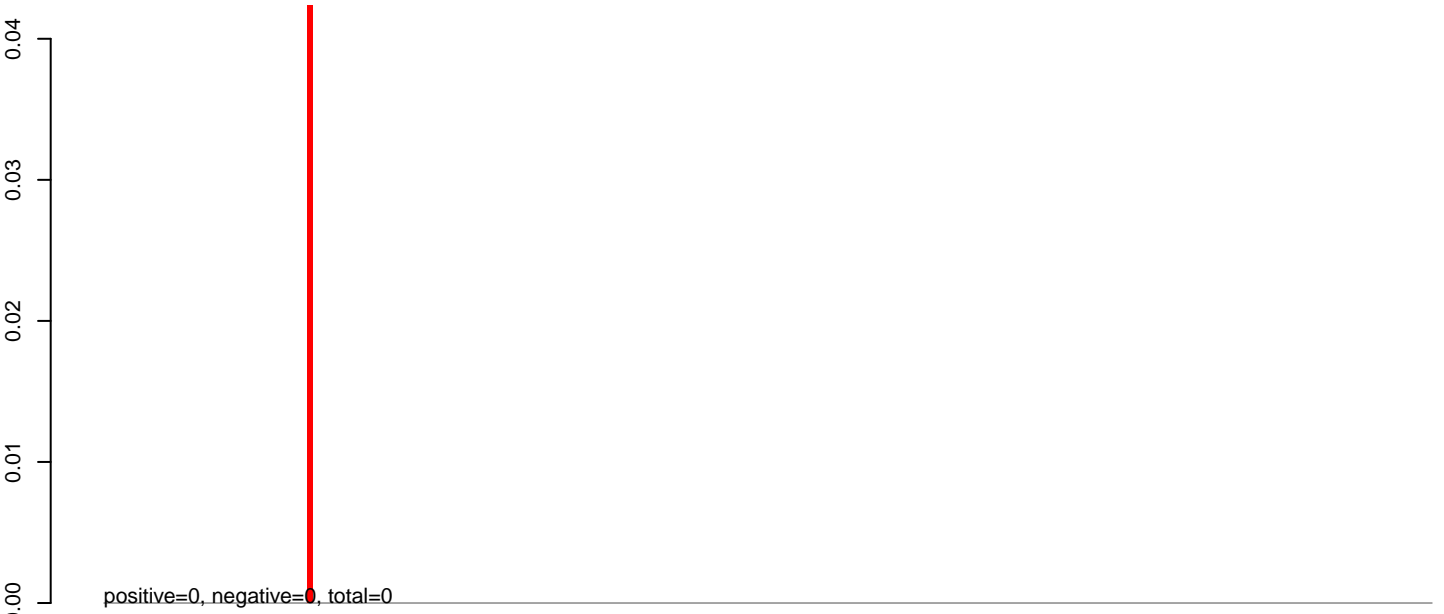
AeAlbo_C636_cells.rep



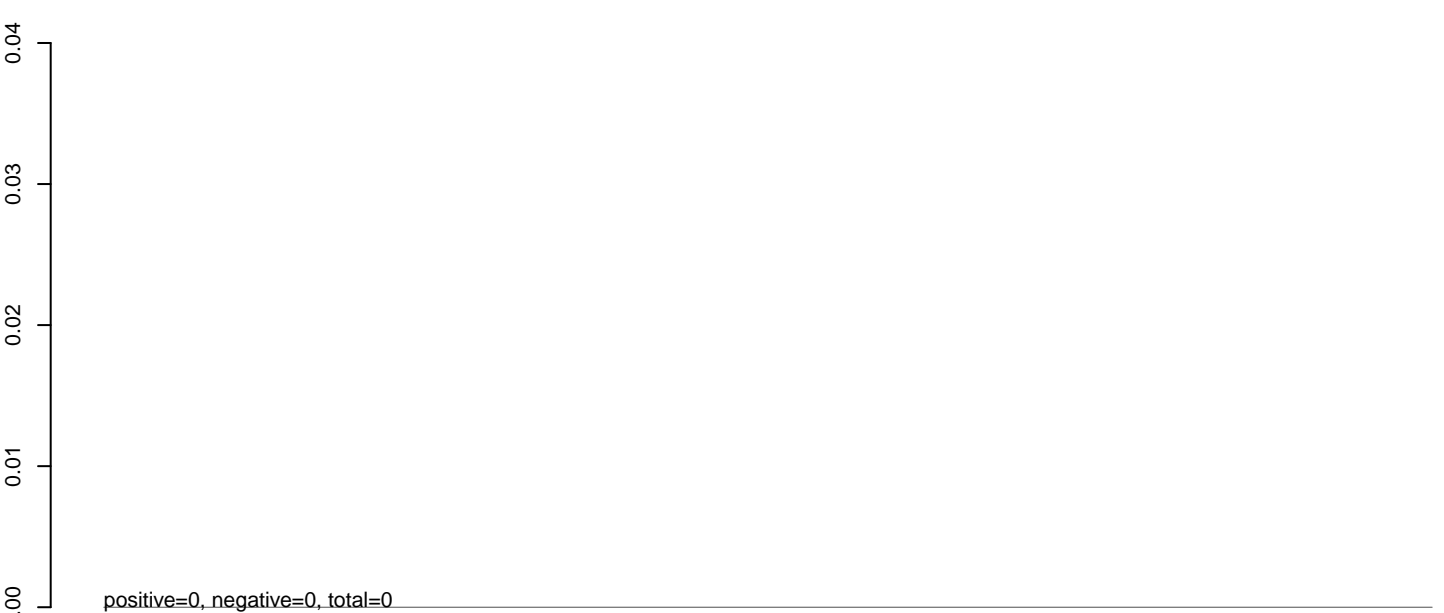
NC_038707.1-Jonchet_virus_strain_B81-CI-2004_segment_S_plus_segment_M_plus_segment_L:1-14098

0 2000 4000 6000 8000 10000 12000 14000

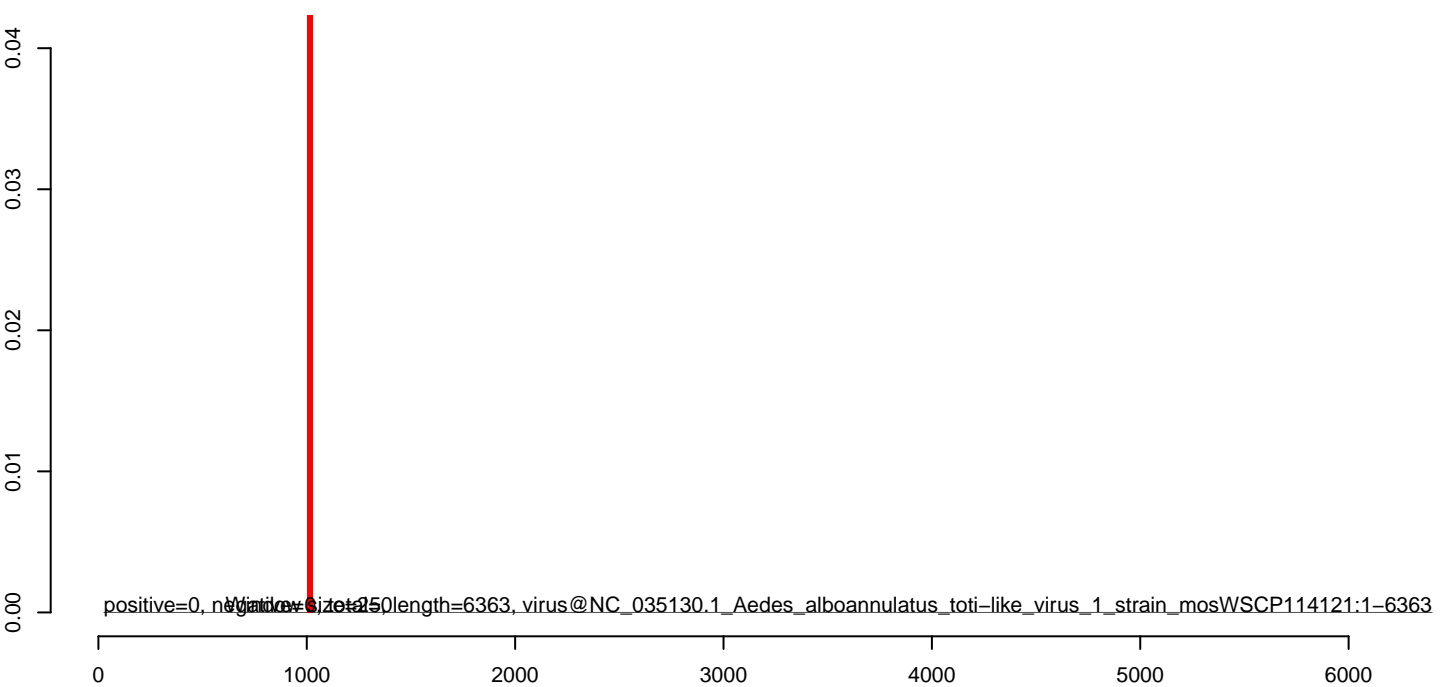
AeAlbo_C636_cells.18_23.rep



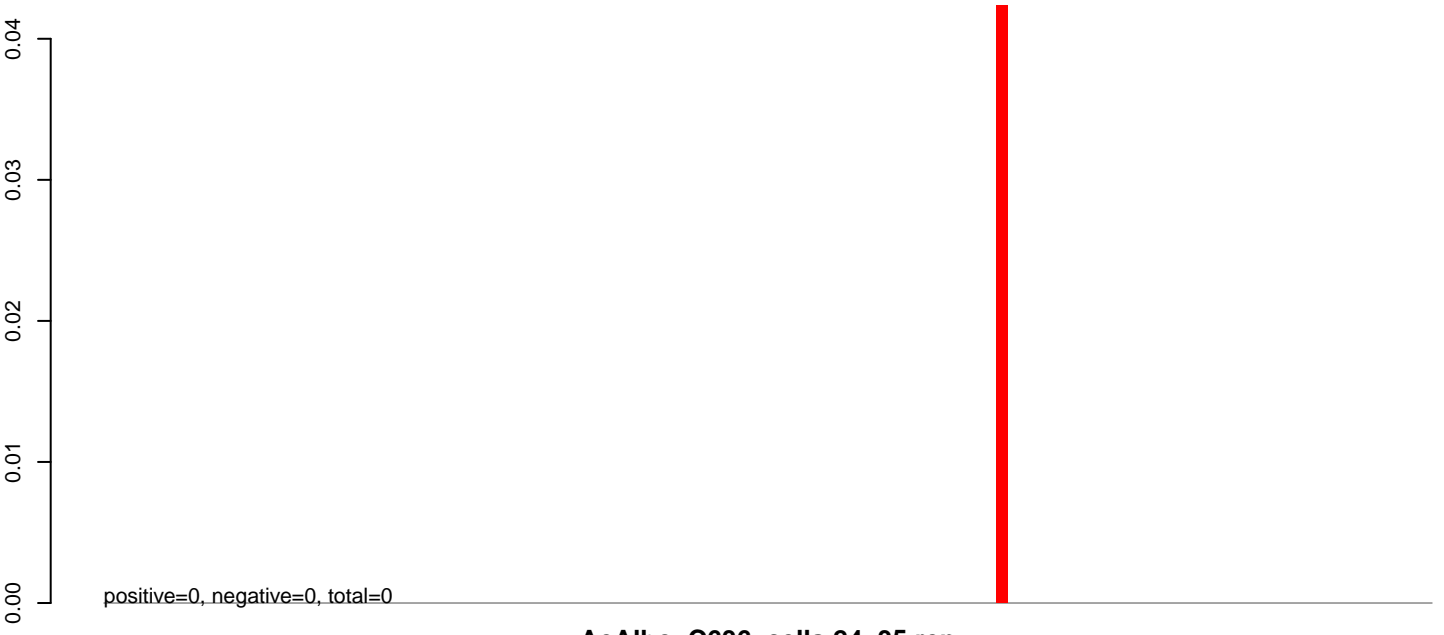
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



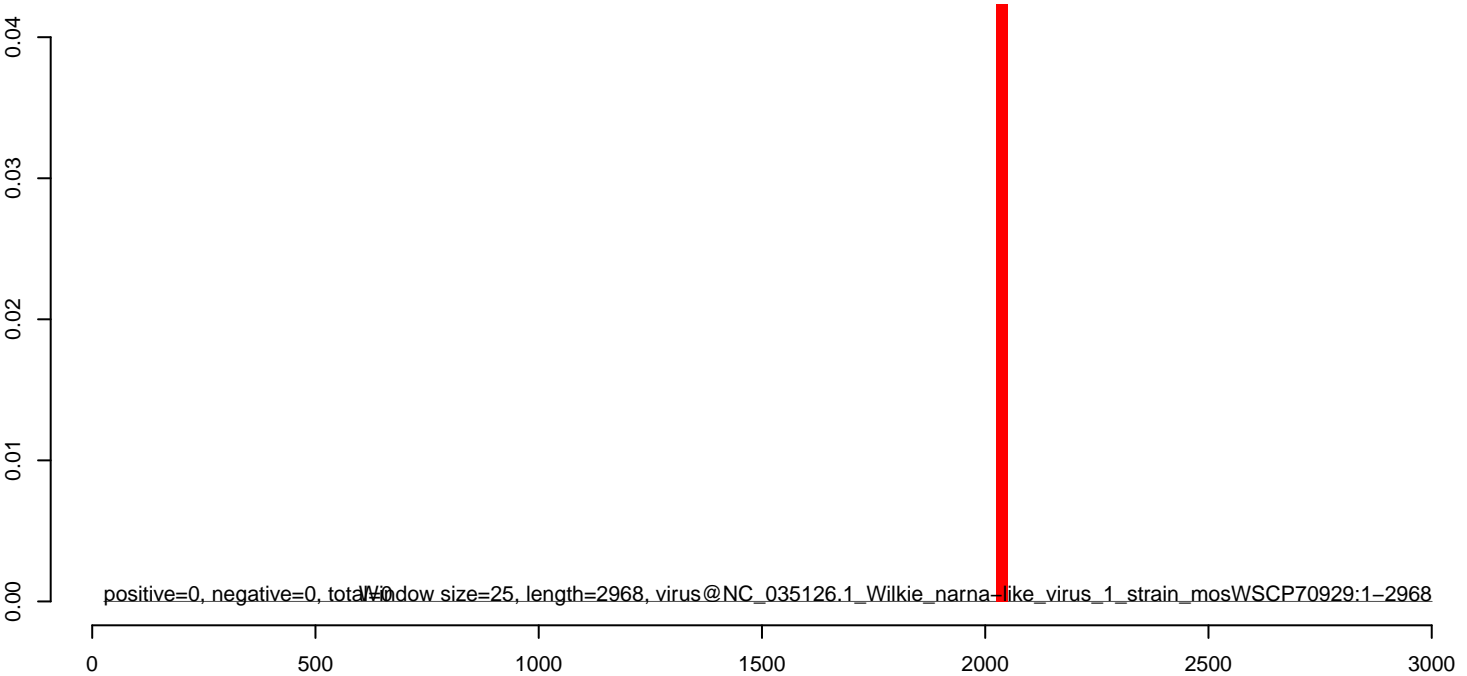
AeAlbo_C636_cells.18_23.rep



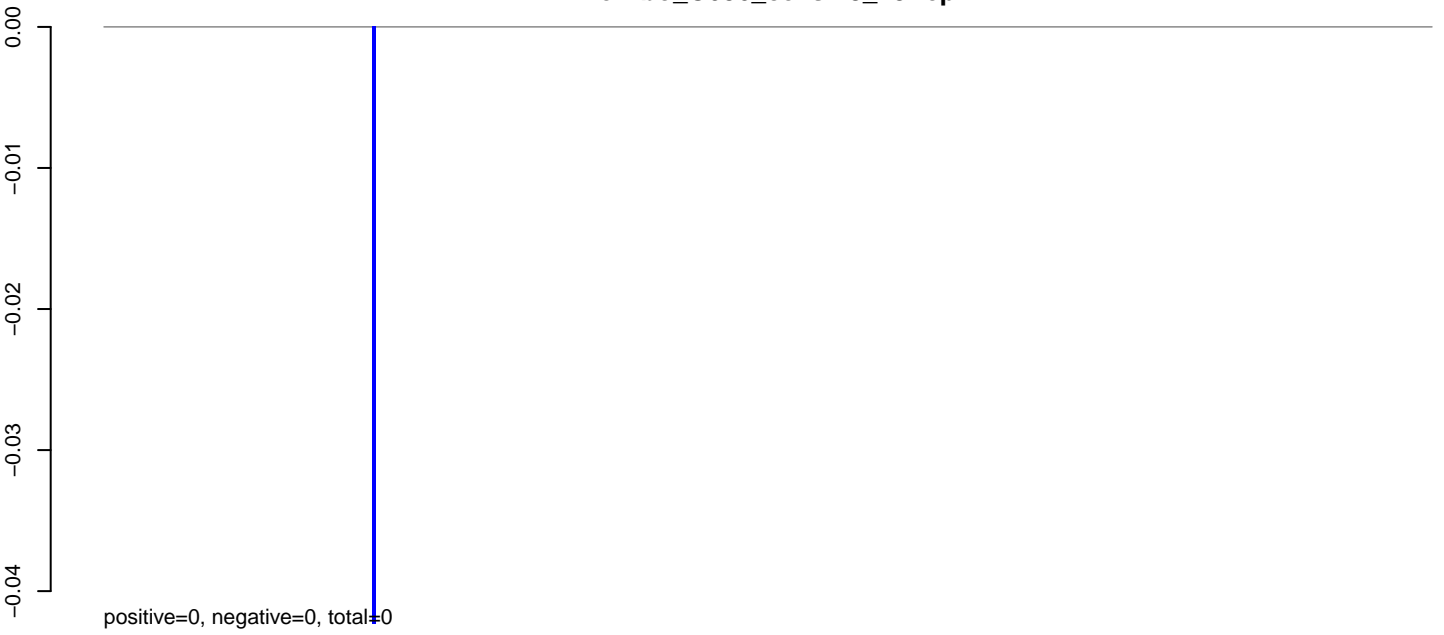
AeAlbo_C636_cells.24_35.rep



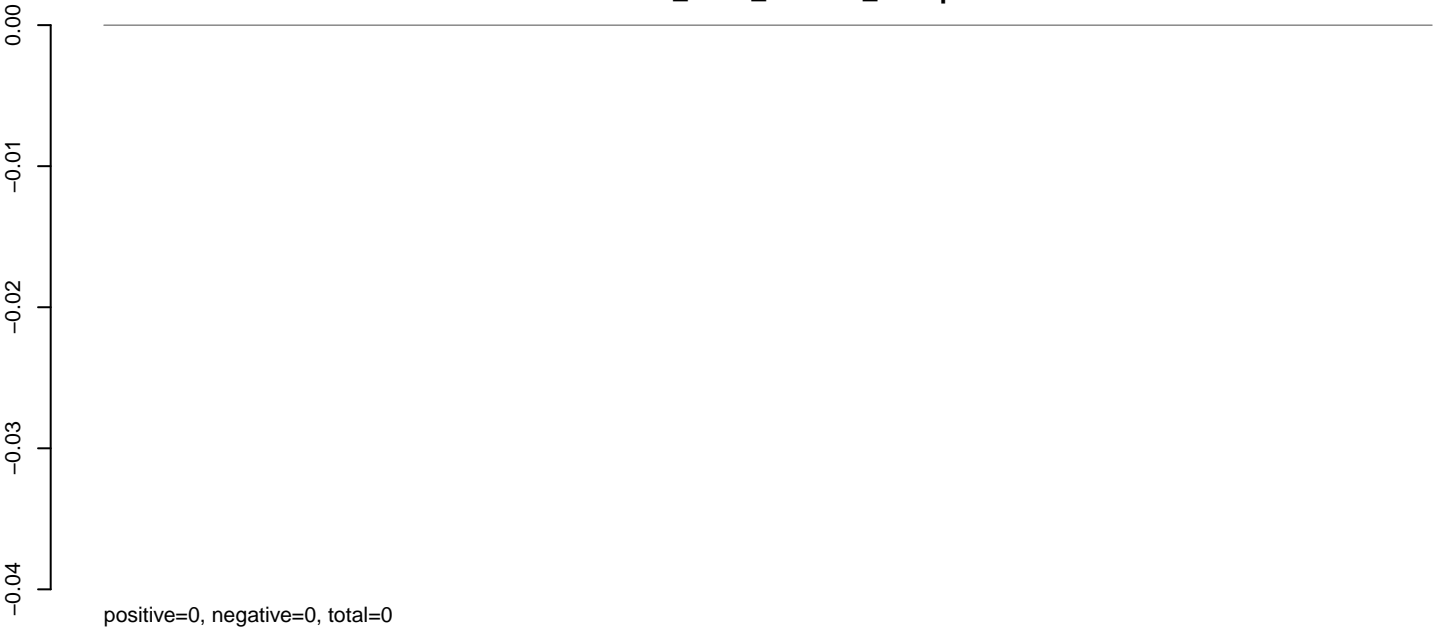
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



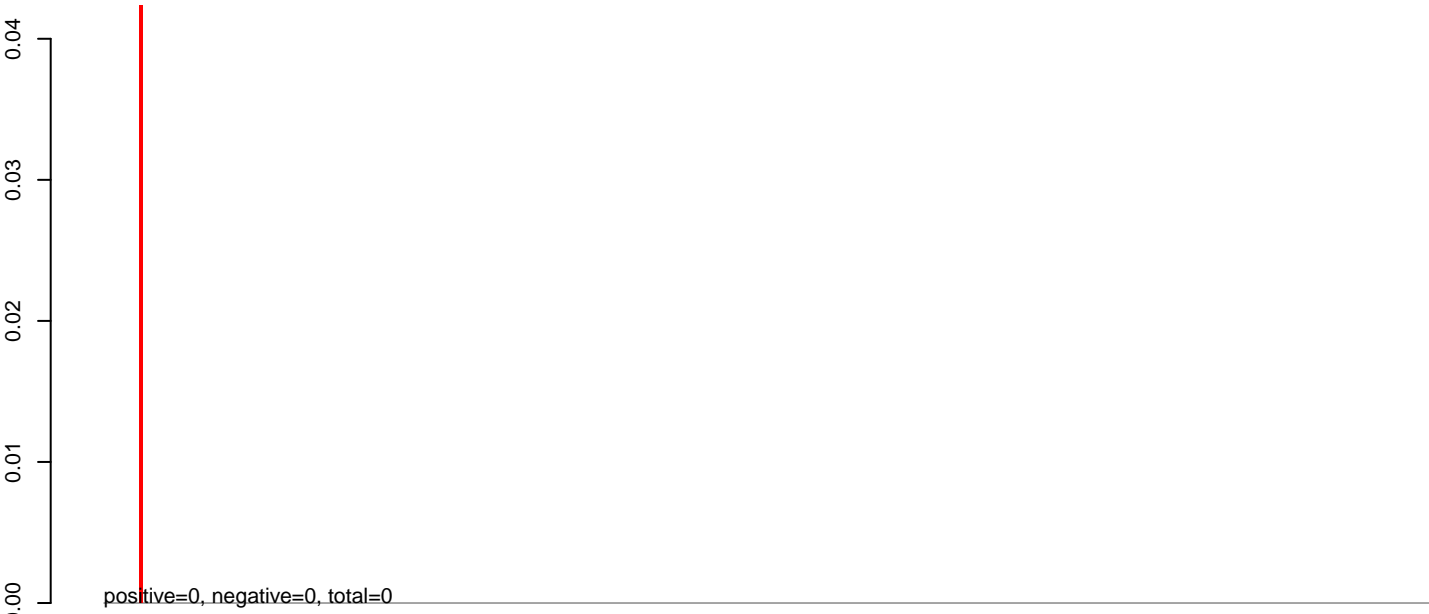
AeAlbo_C636_cells.rep



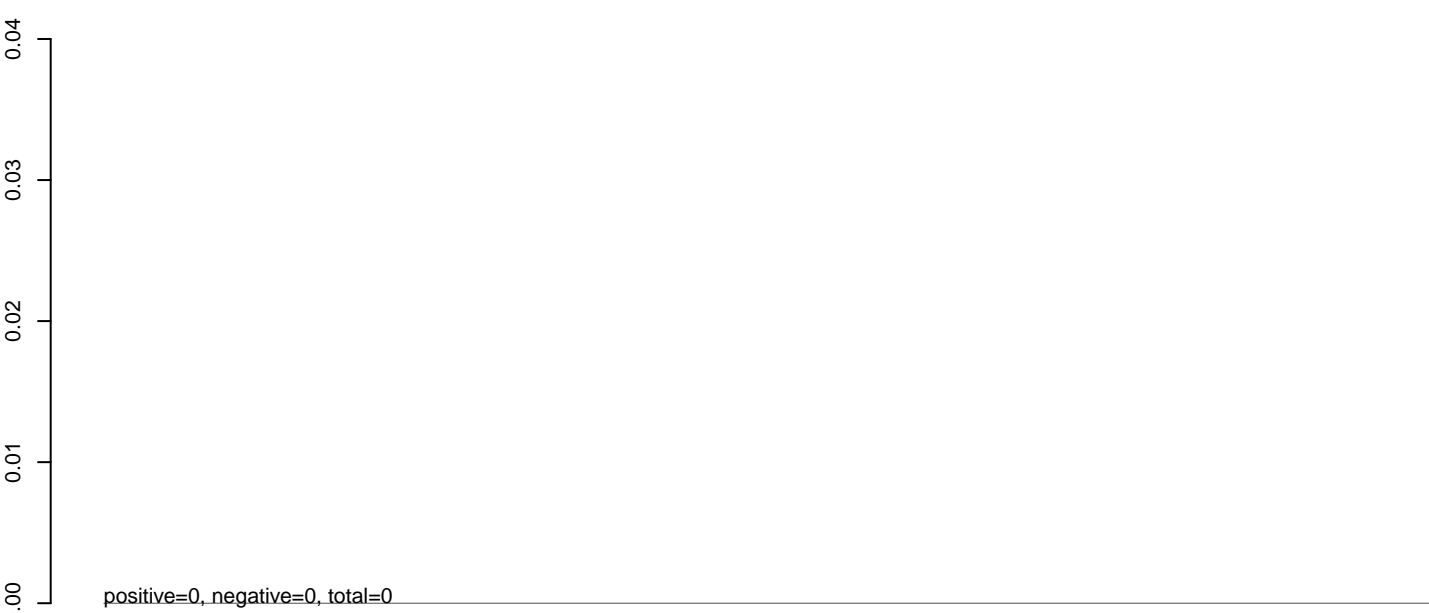
bow size style=0, the=0, int=0 NC_034454.1-Fukuoka_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_protein_glycoprotei:1-10863

0 2000 4000 6000 8000 10000

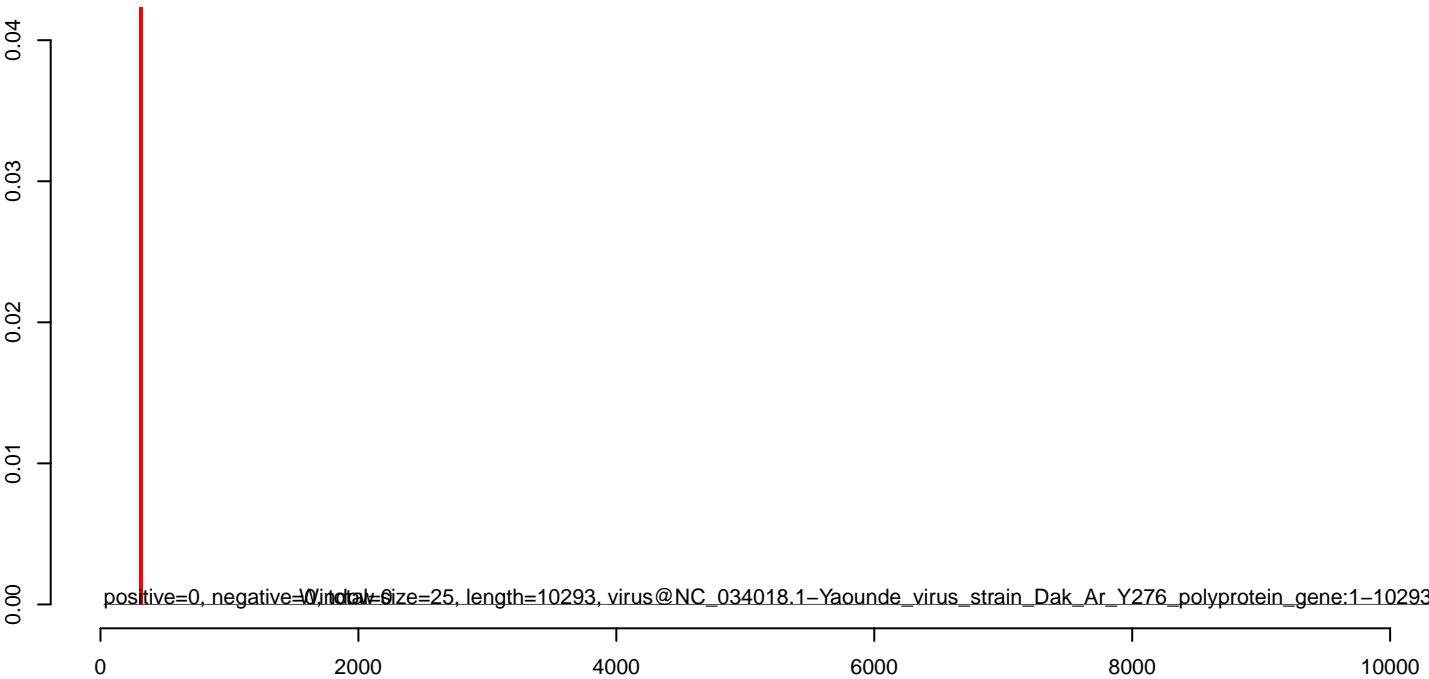
AeAlbo_C636_cells.18_23.rep



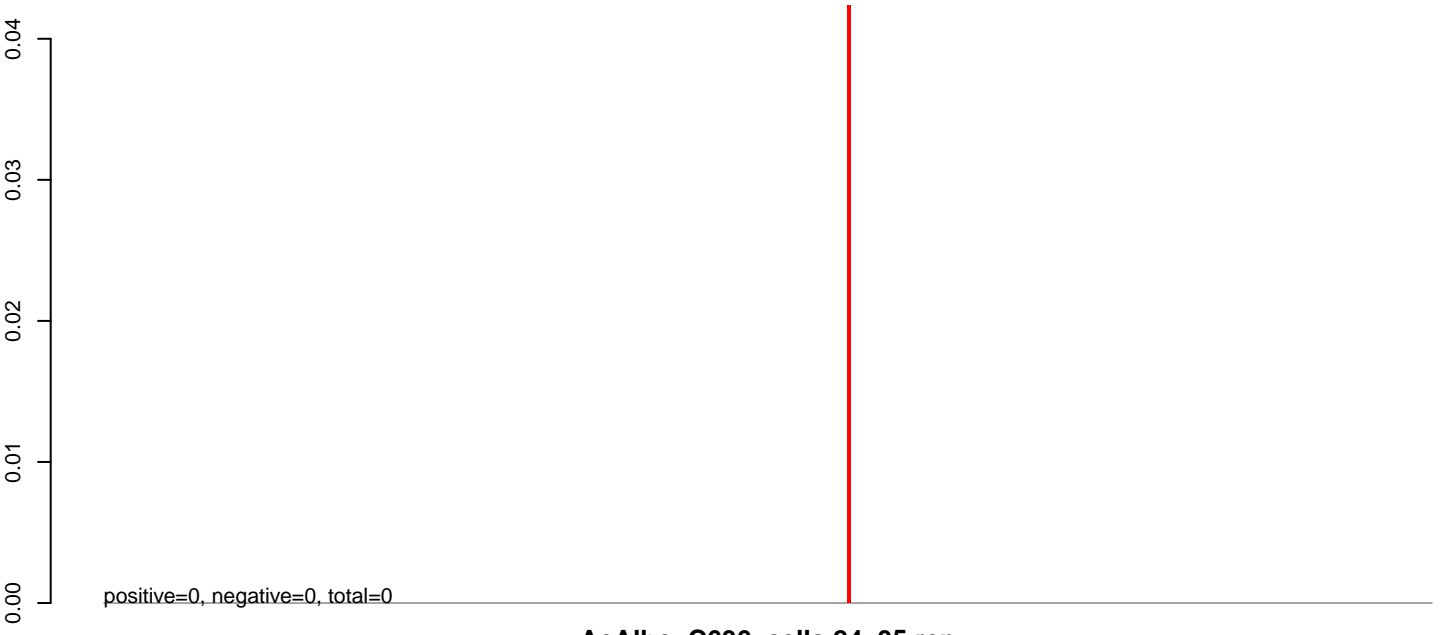
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



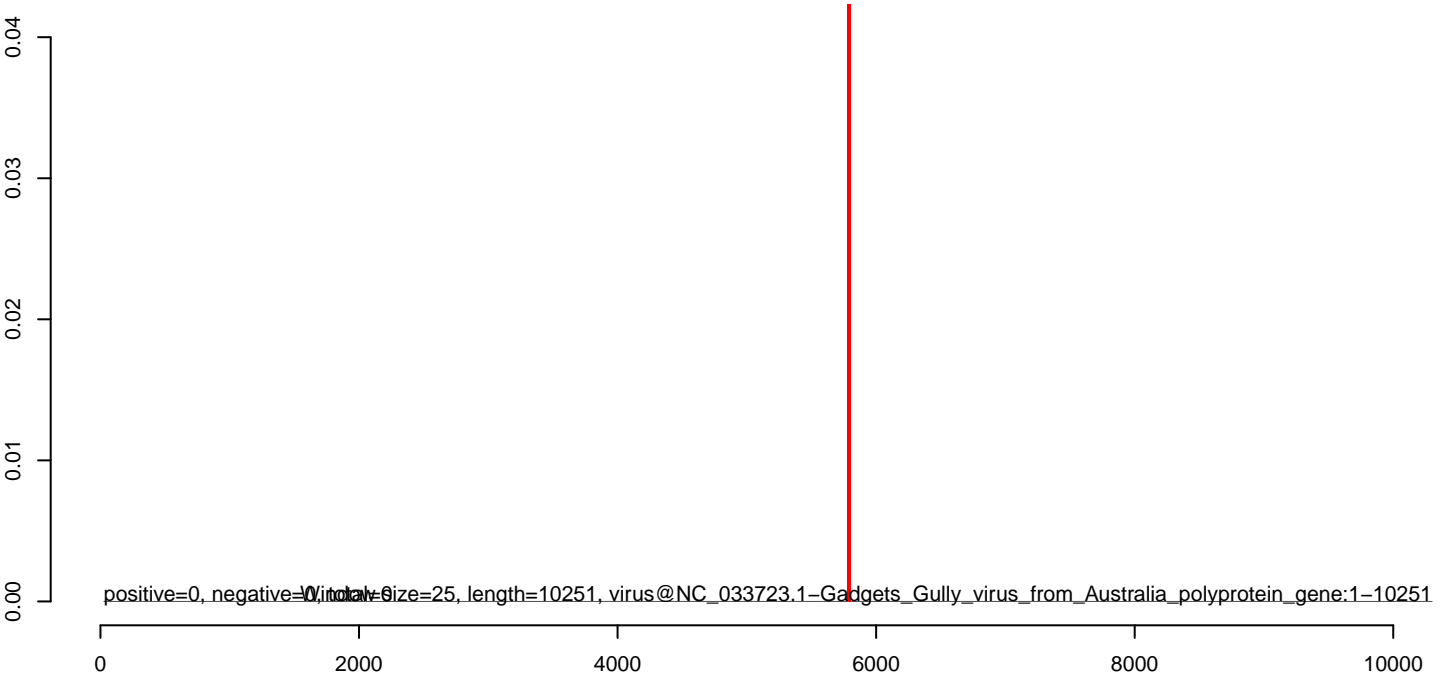
AeAlbo_C636_cells.18_23.rep



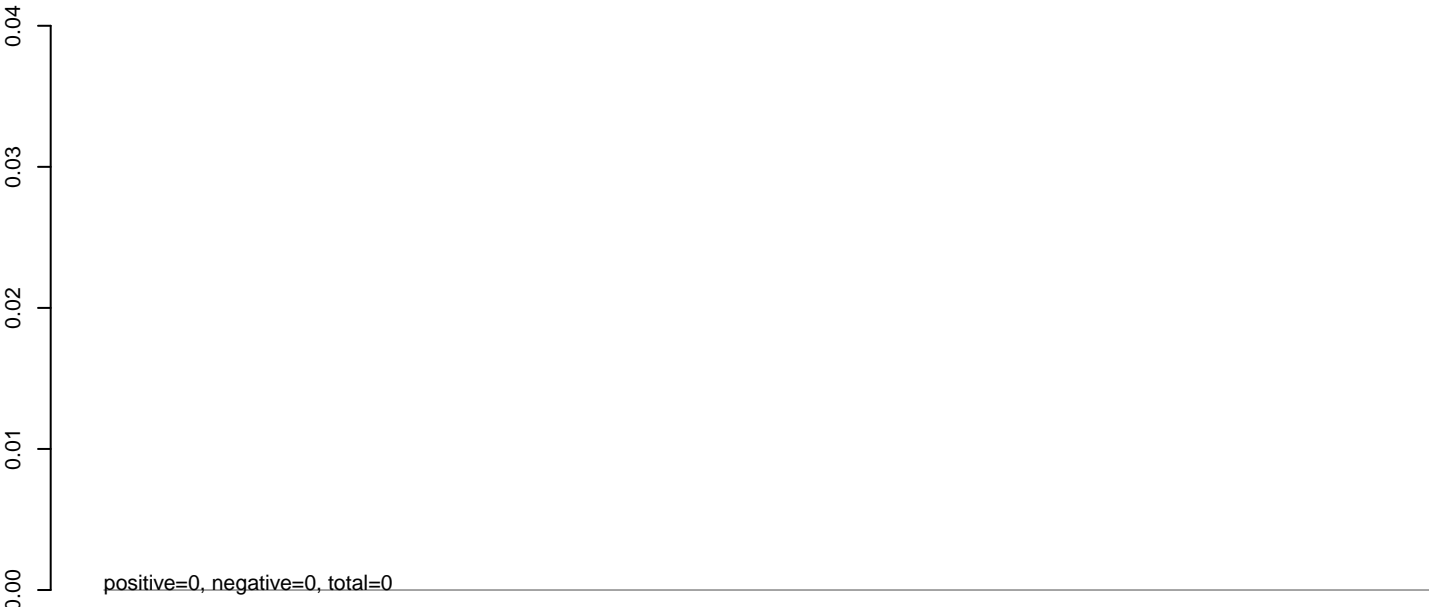
AeAlbo_C636_cells.24_35.rep



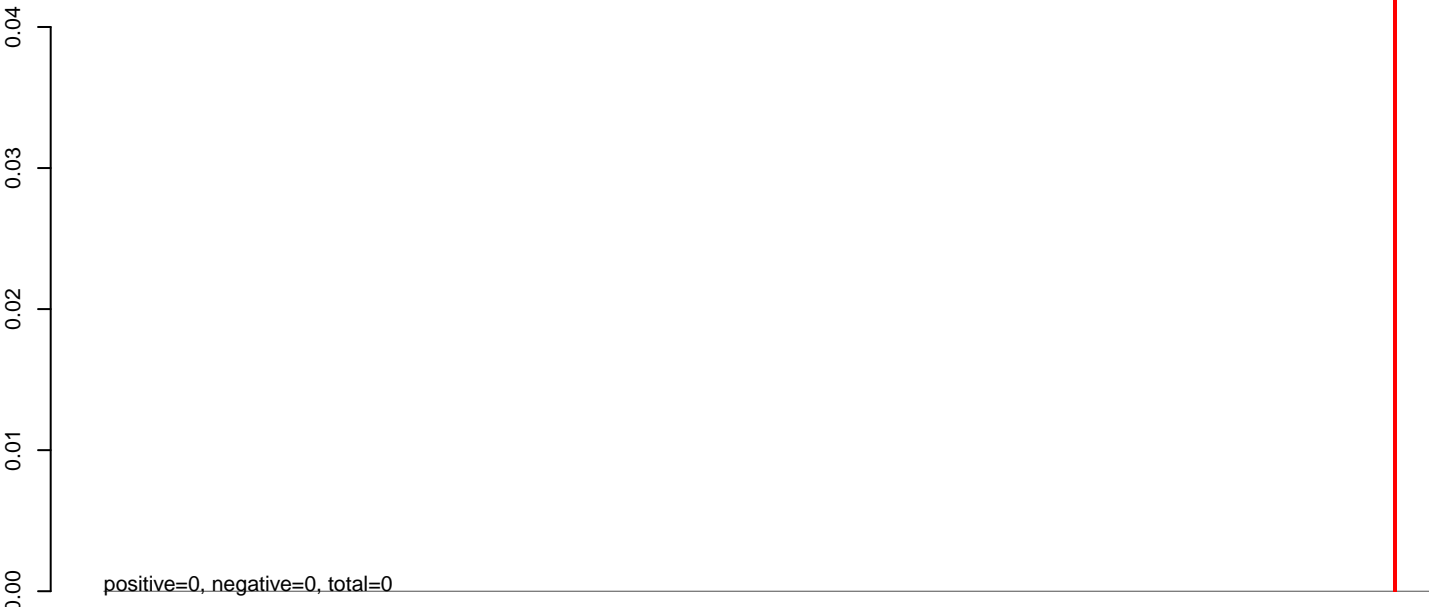
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

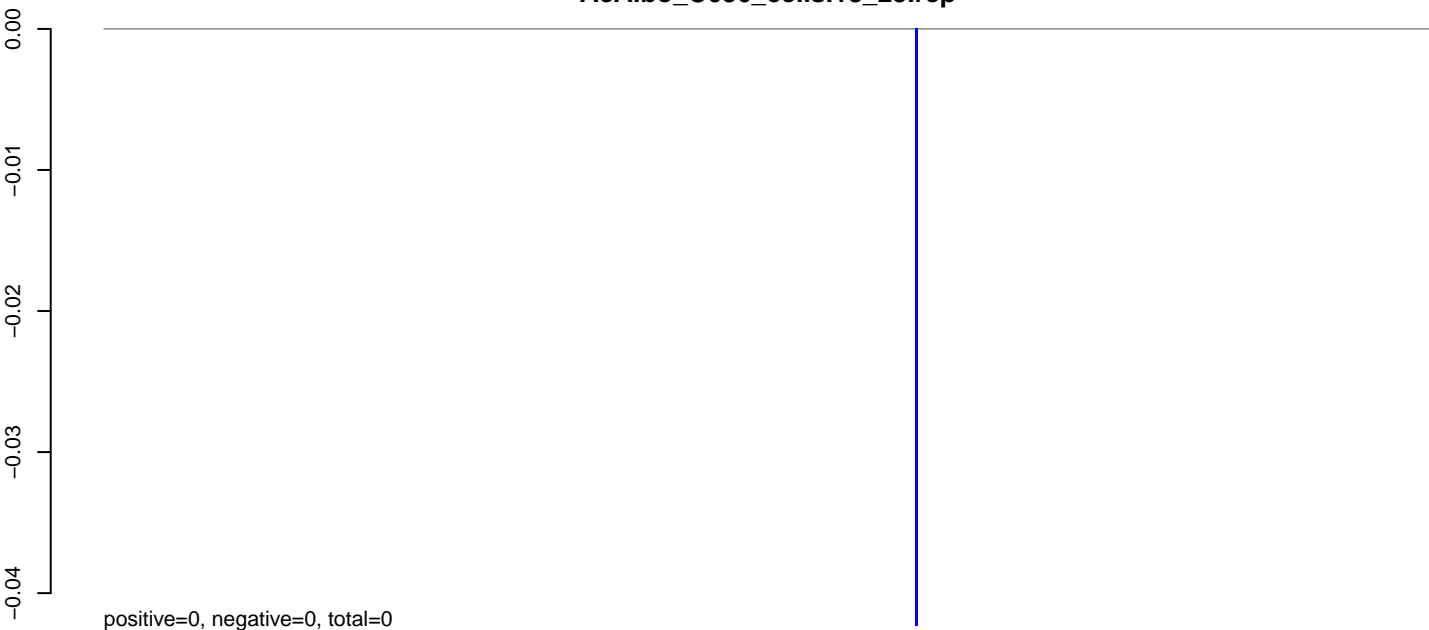


AeAlbo_C636_cells.rep

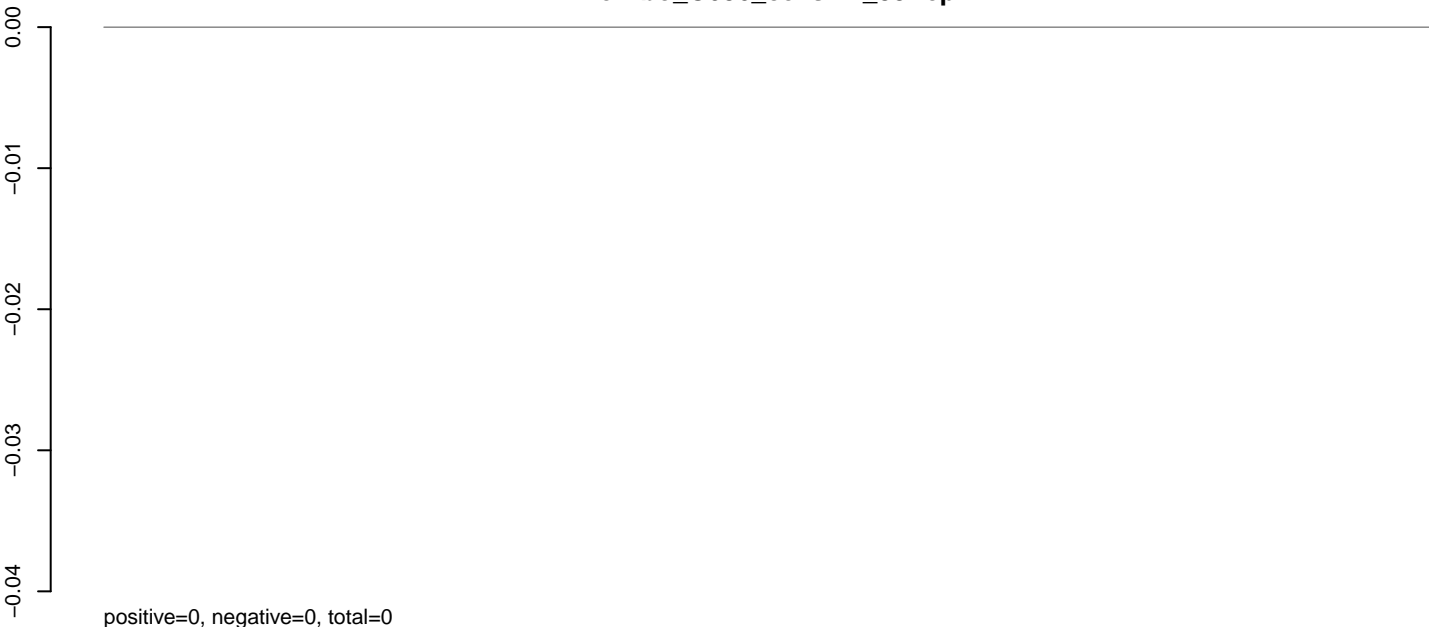


0 2000 4000 6000 8000 10000

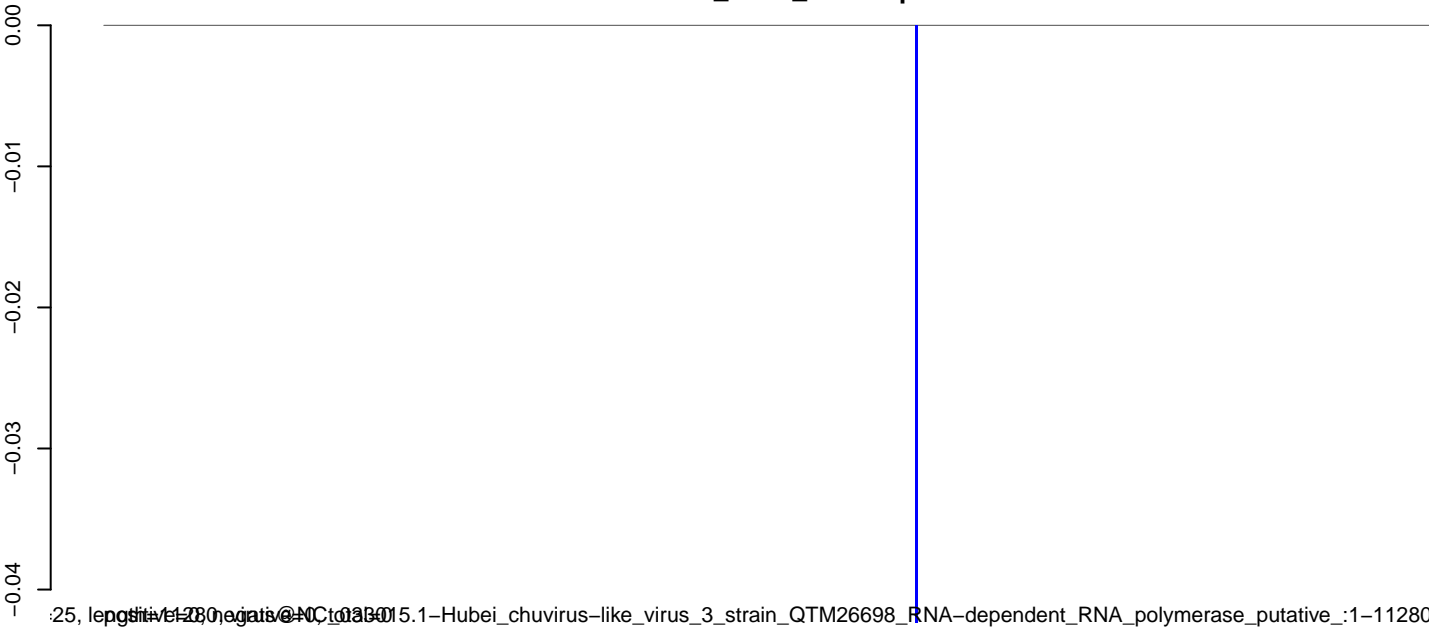
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



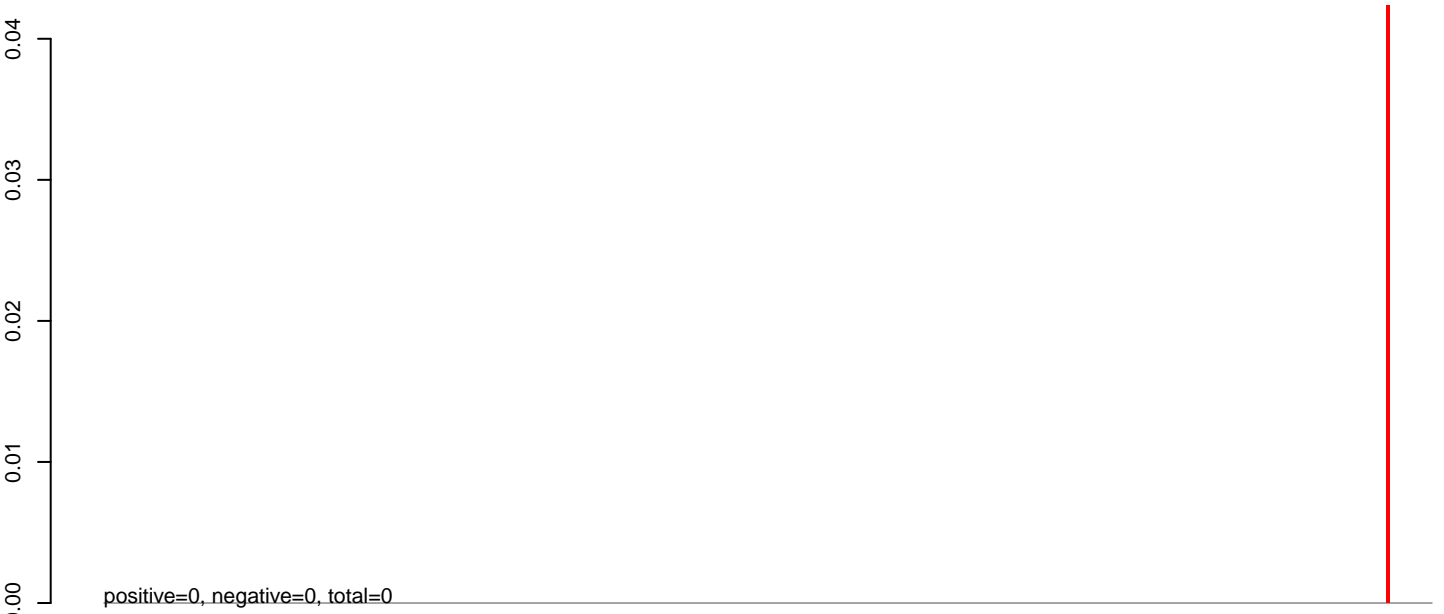
AeAlbo_C636_cells.rep



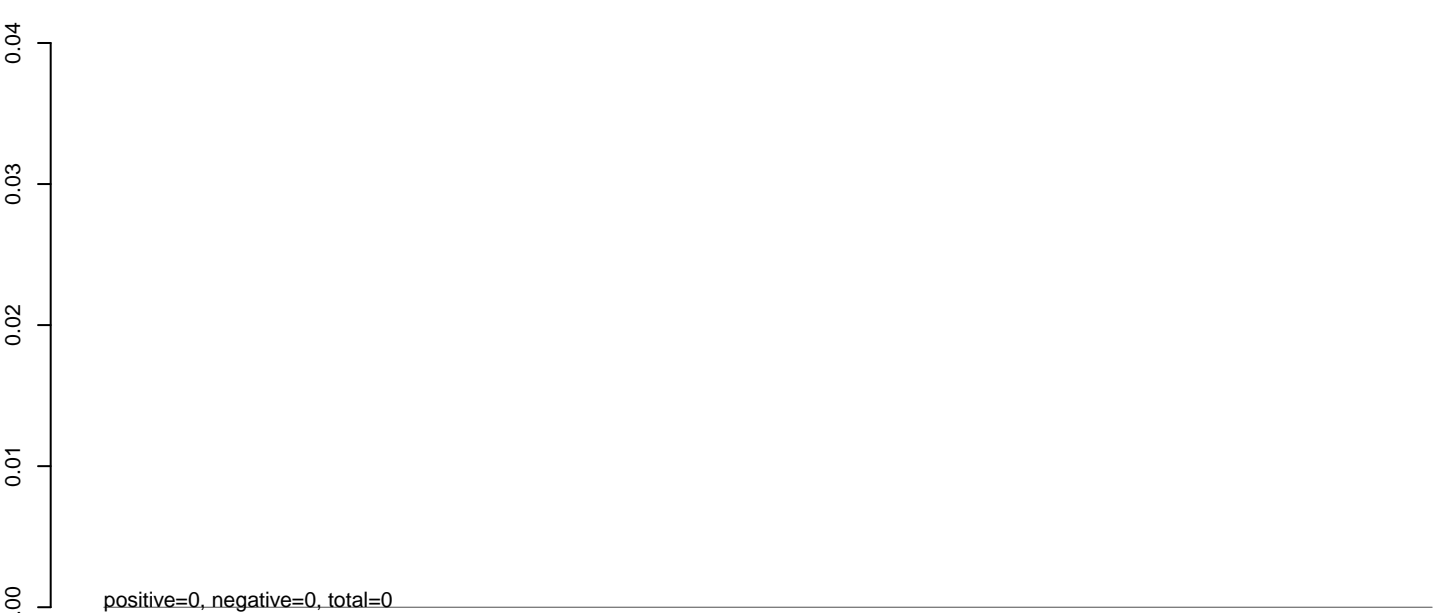
25, length=280, negative=0, total=0 15.1-Hubei_chuvirus-like_virus_3_strain_QTM26698_RNA-dependent_RNA_polymerase_putative_:1-11280

0 2000 4000 6000 8000 10000

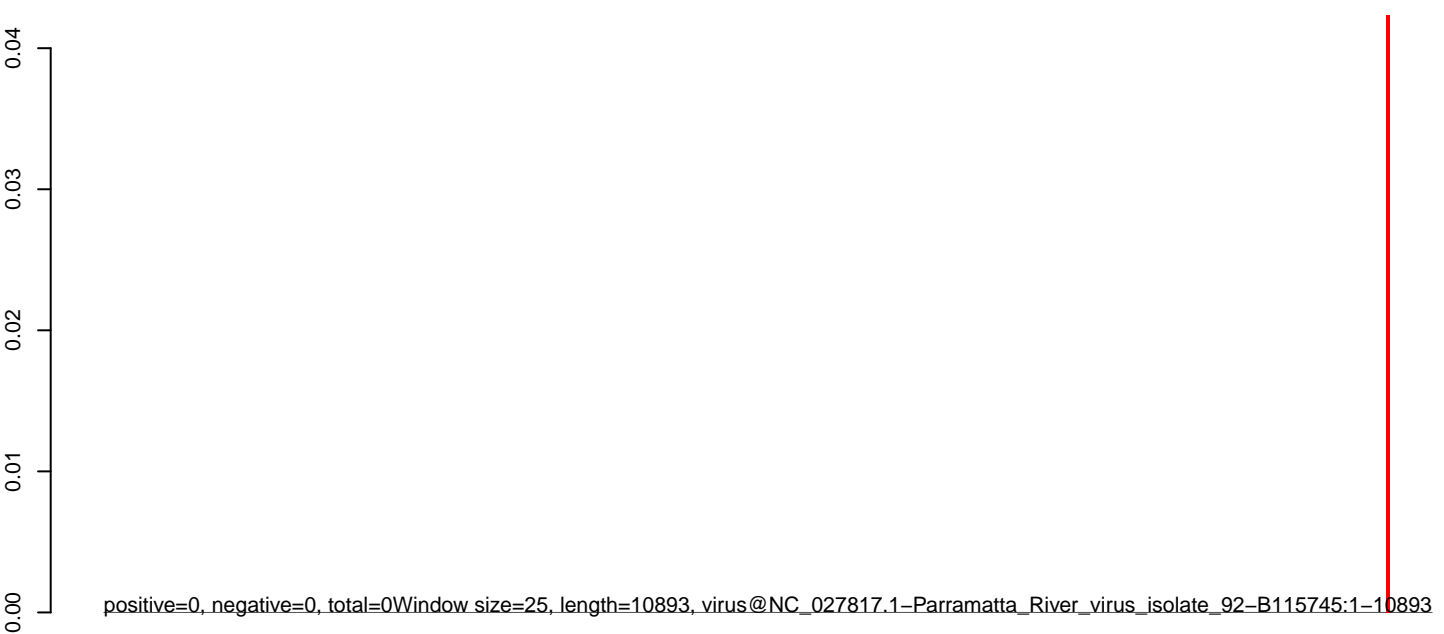
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



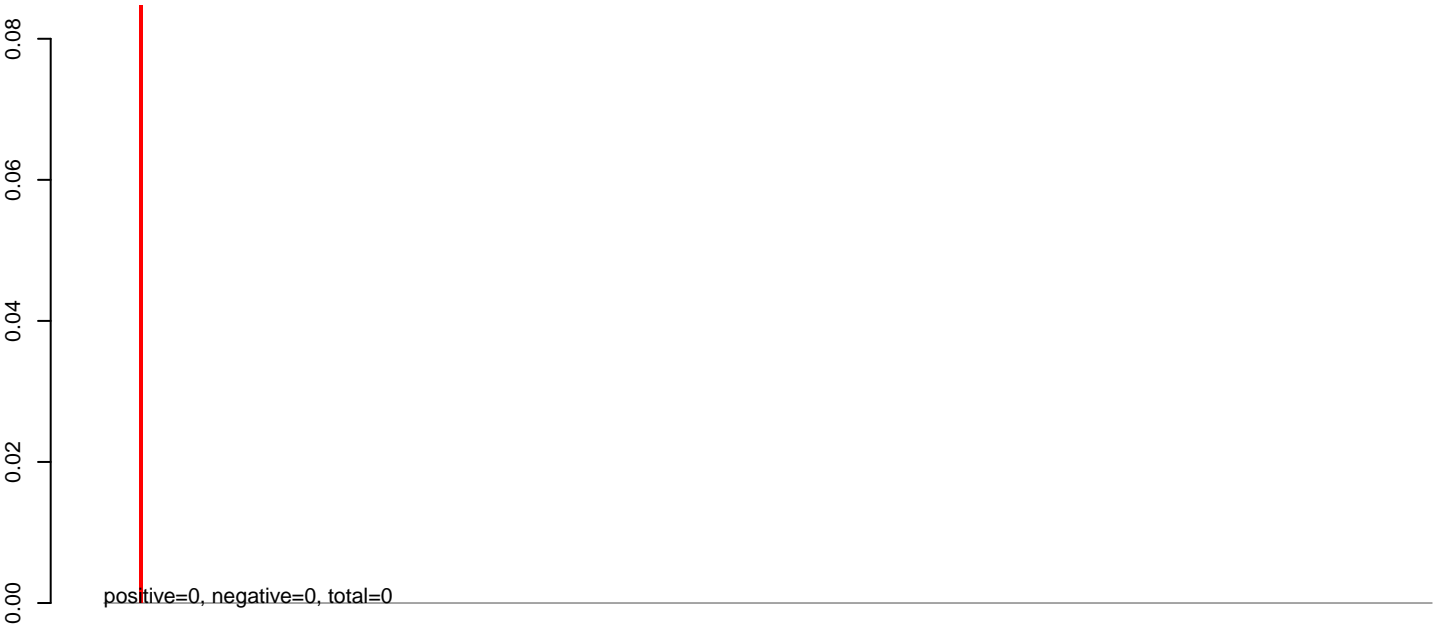
AeAlbo_C636_cells.rep



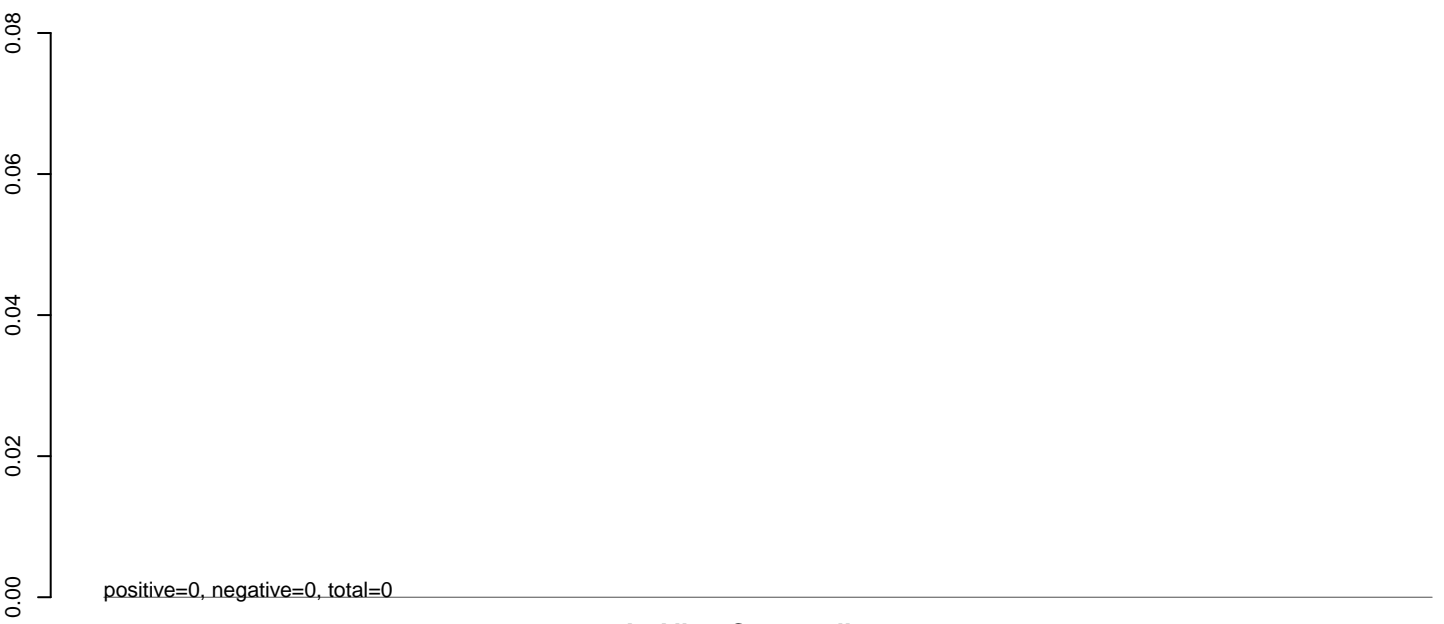
Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



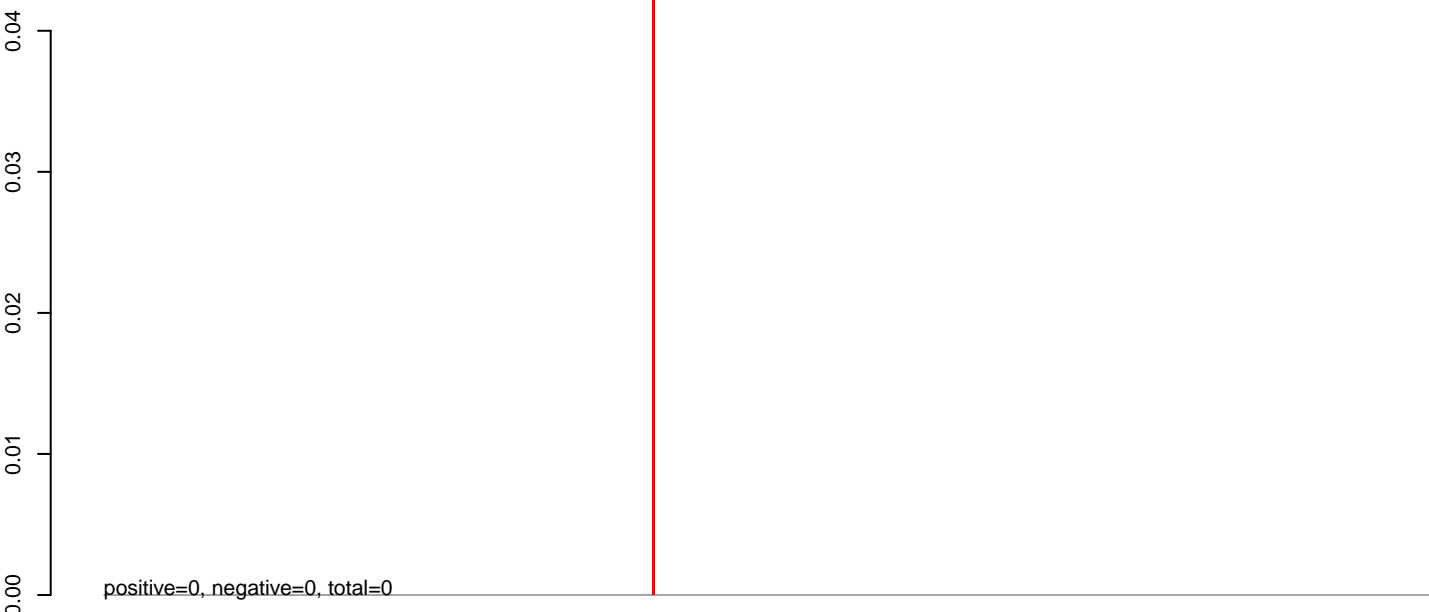
AeAlbo_C636_cells.24_35.rep



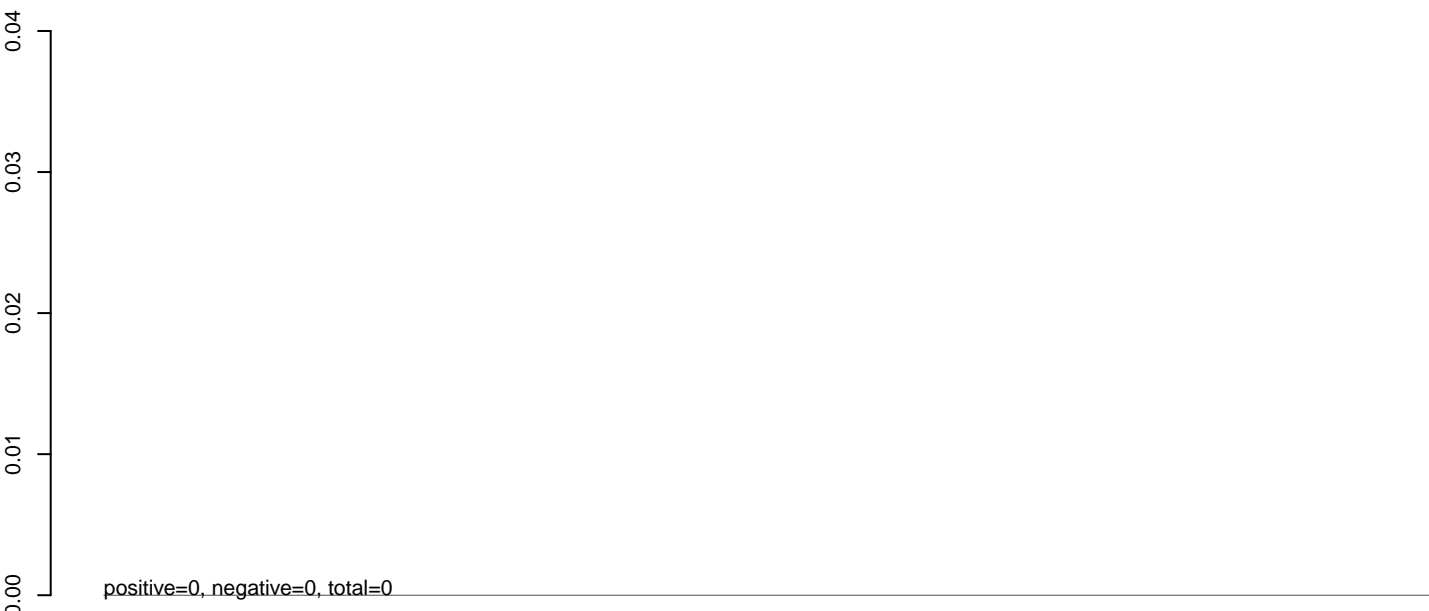
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

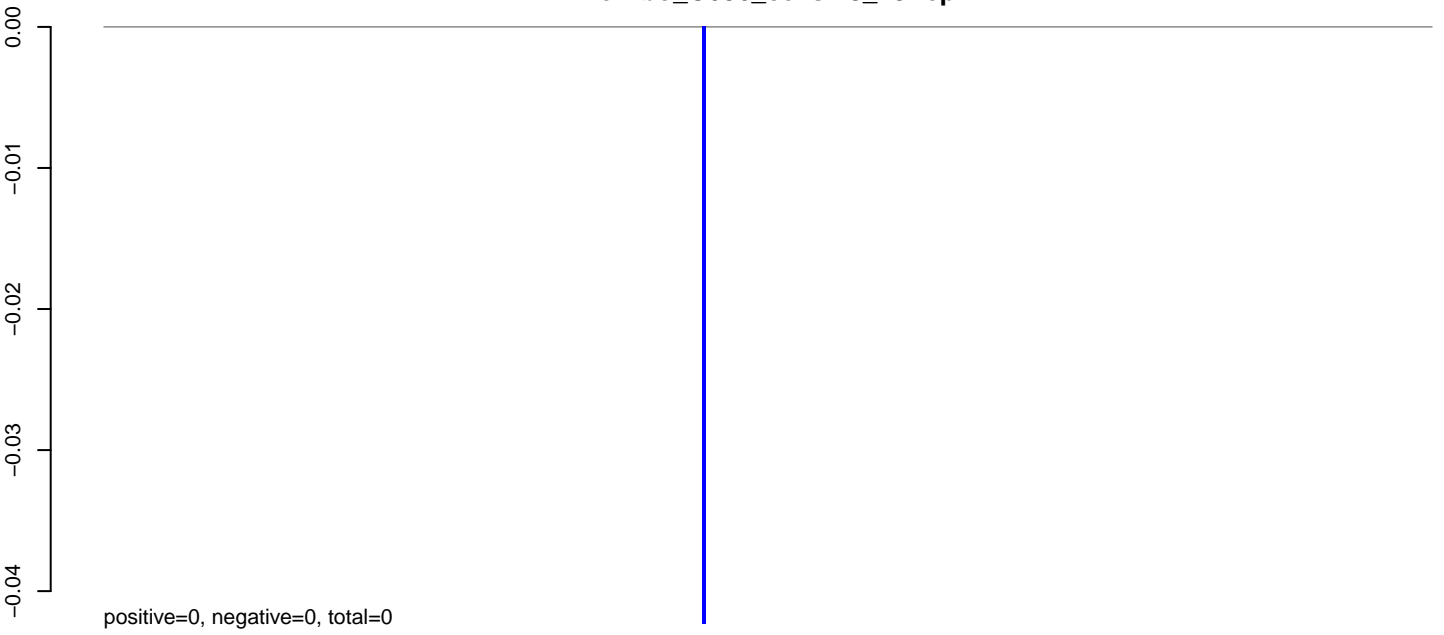


AeAlbo_C636_cells.rep

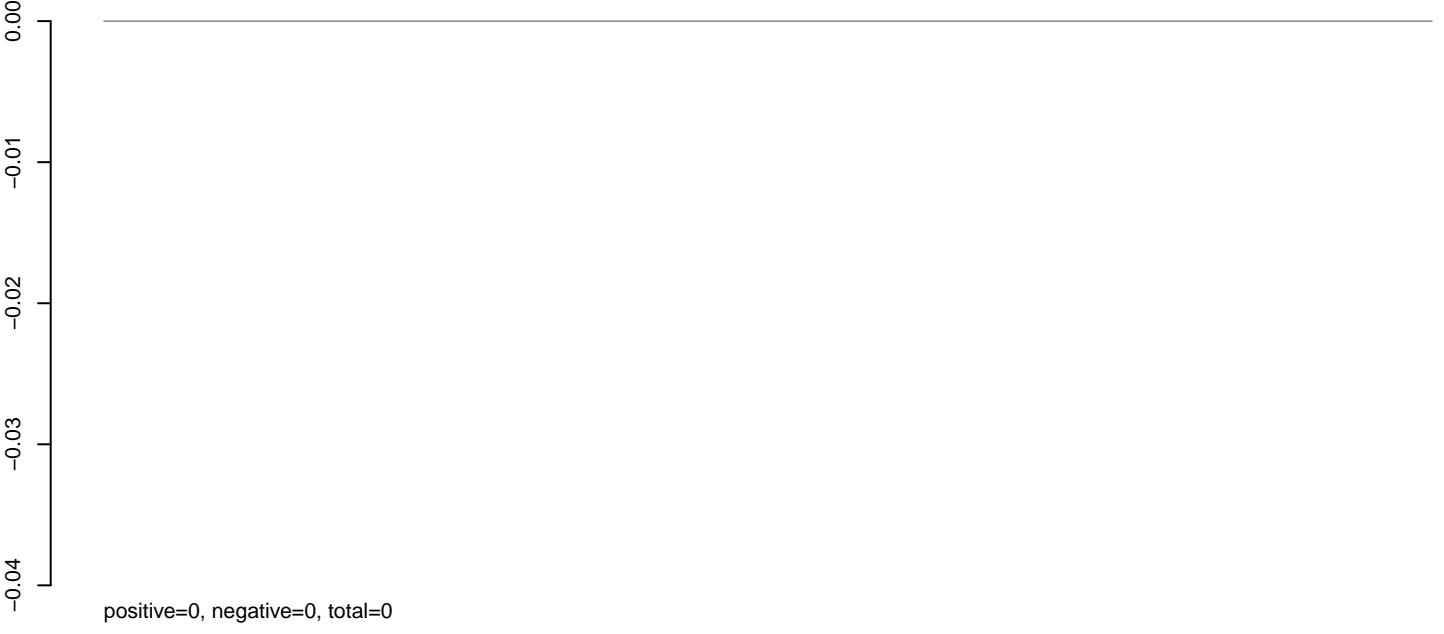


Window size=25, length=11190, virus@NC_025384.1-Culex tritaeniorhynchus rhabdovirus RNA strain_TY:1-11190

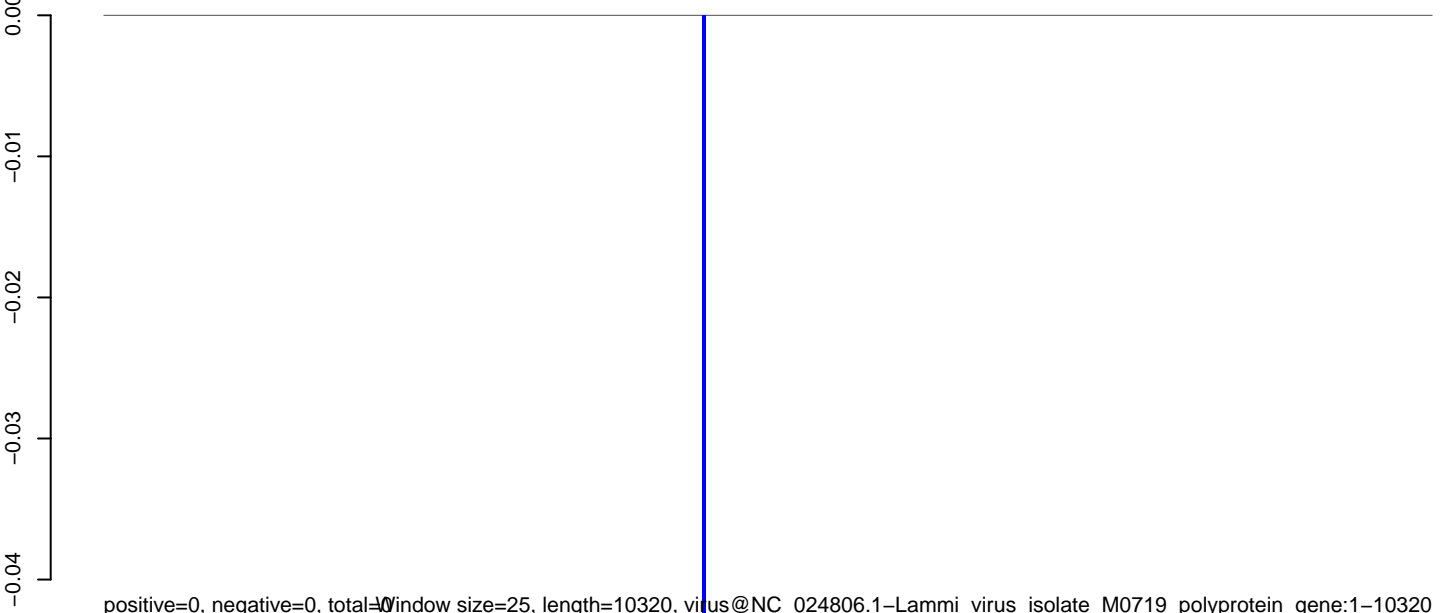
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



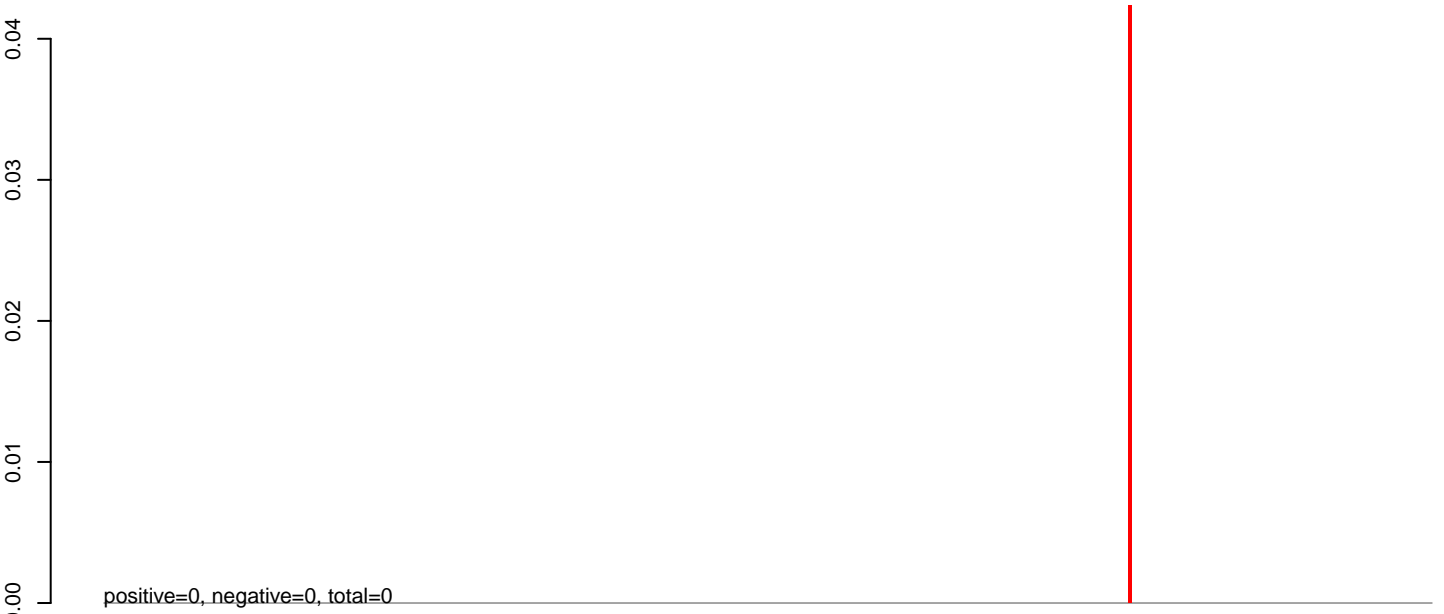
AeAlbo_C636_cells.rep



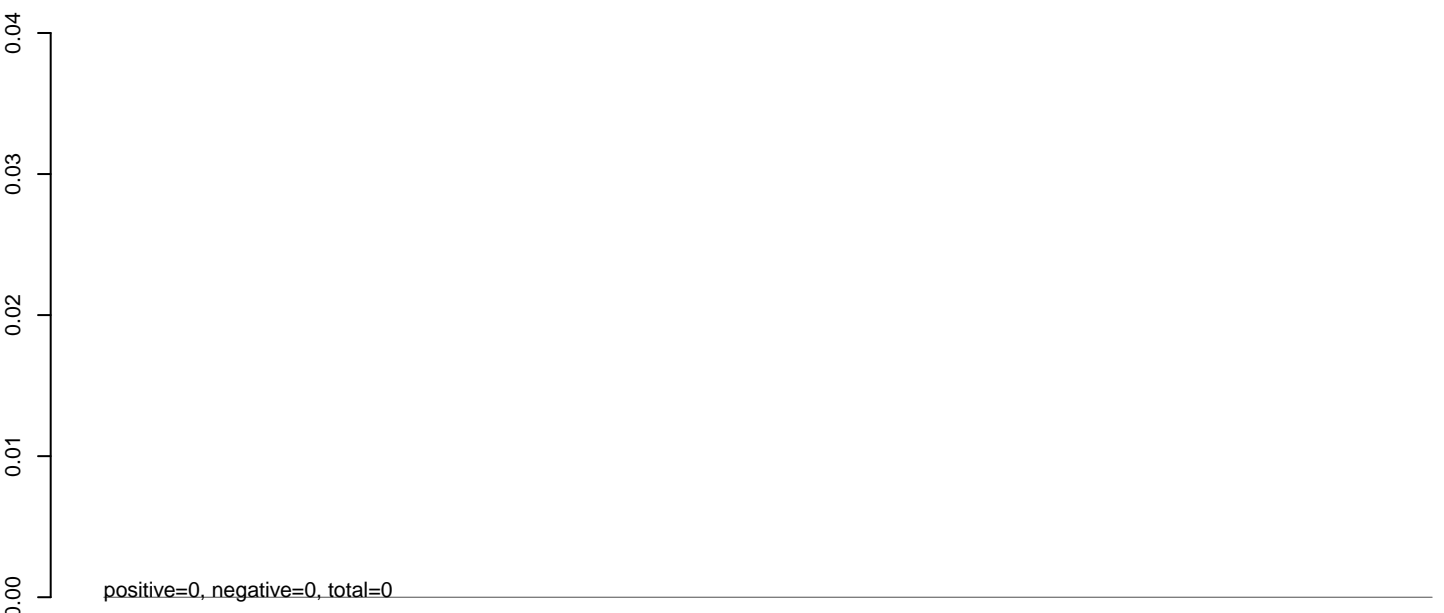
Window size=25, length=10320, virus@NC_024806.1-Lammi_virus_isolate_M0719_polyprotein_gene:1-10320

0 2000 4000 6000 8000 10000

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

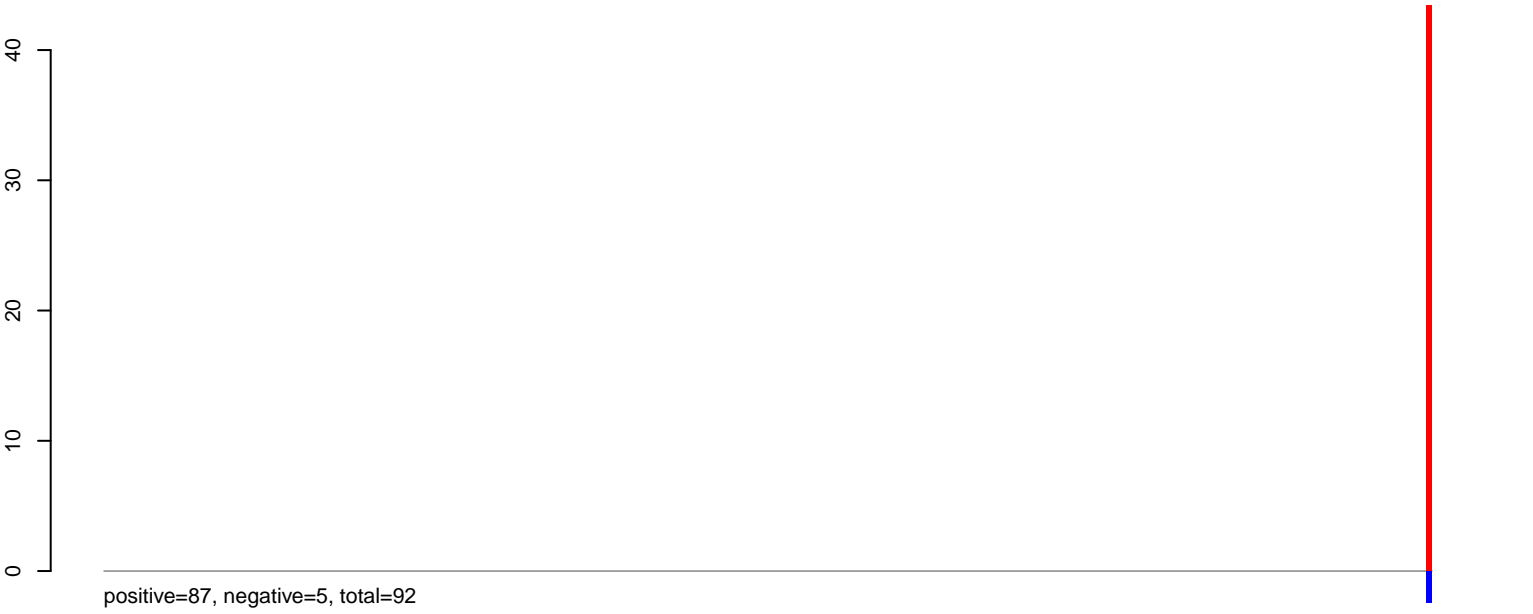


AeAlbo_C636_cells.rep

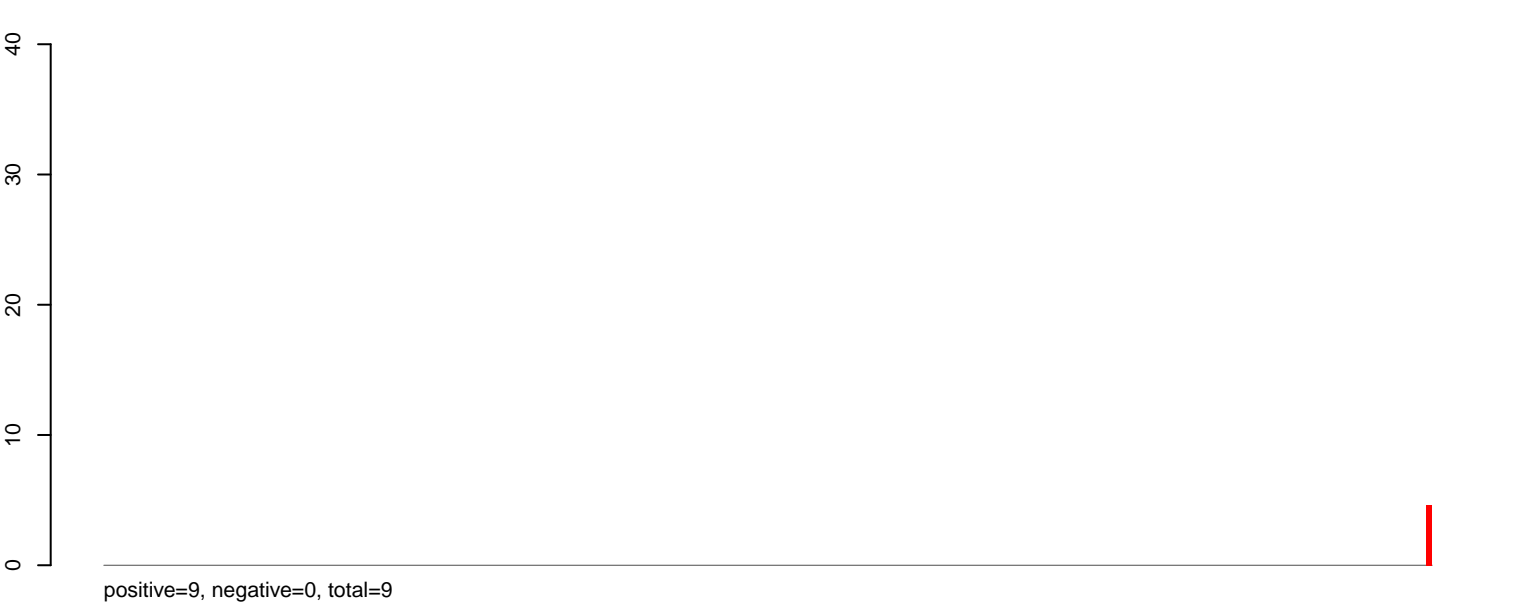


0 2000 4000 6000 8000 10000

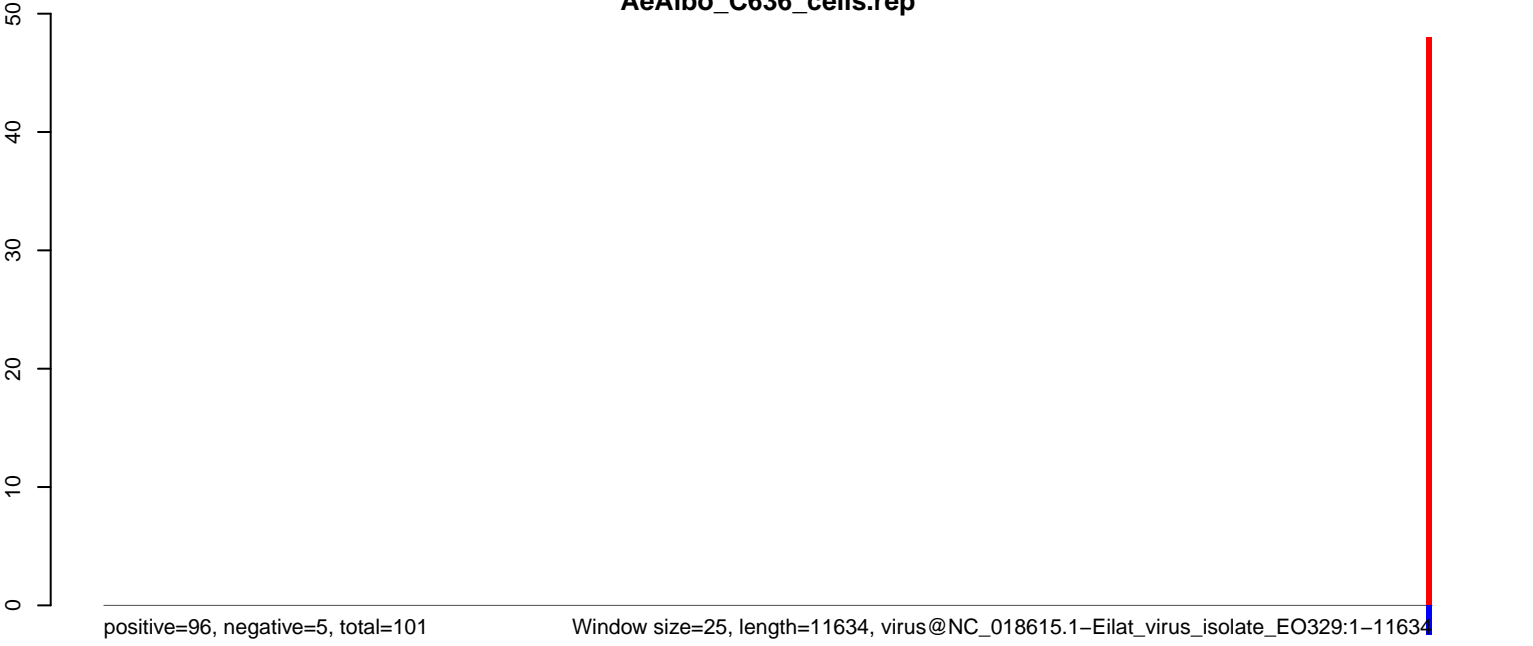
AeAlbo_C636_cells.18_23.rep



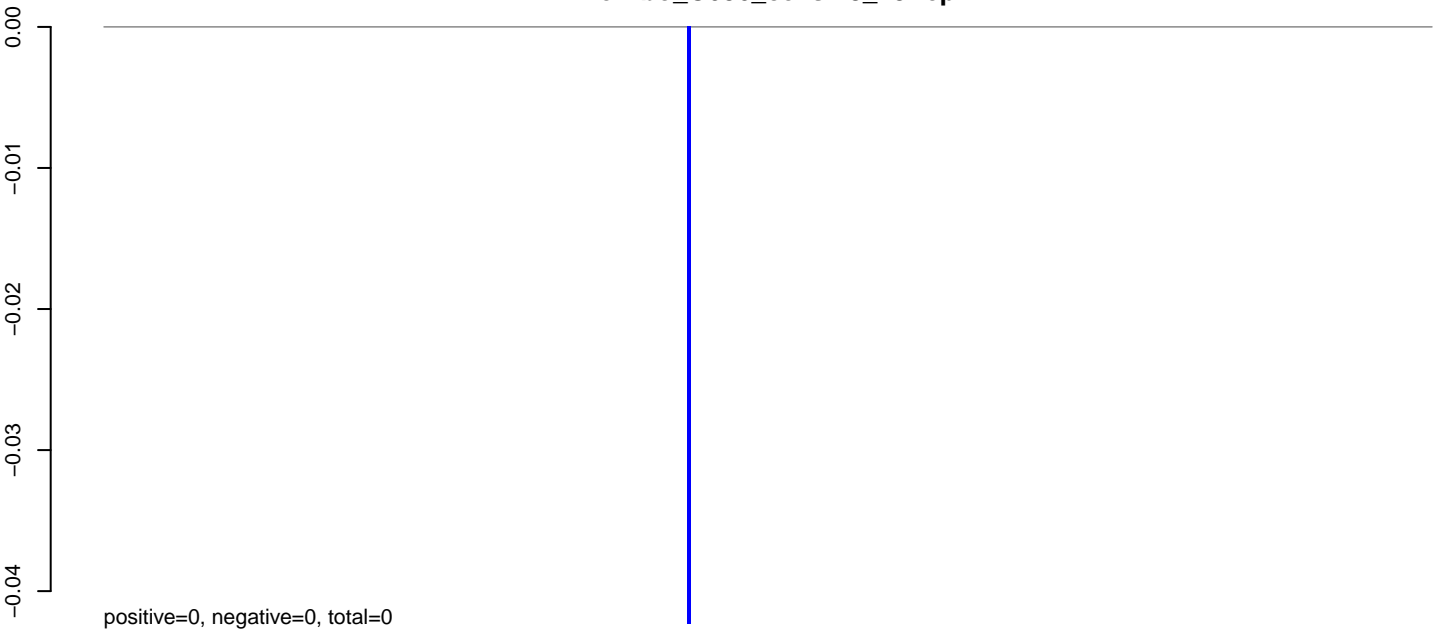
AeAlbo_C636_cells.24_35.rep



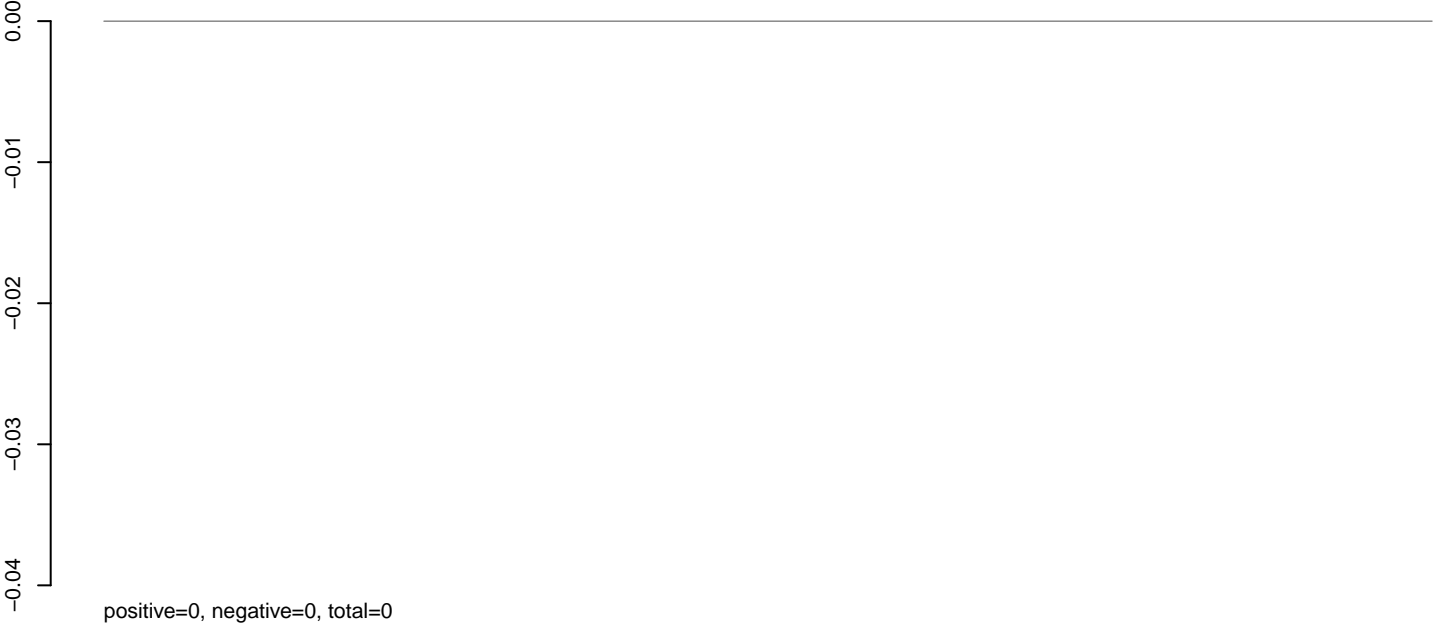
AeAlbo_C636_cells.rep



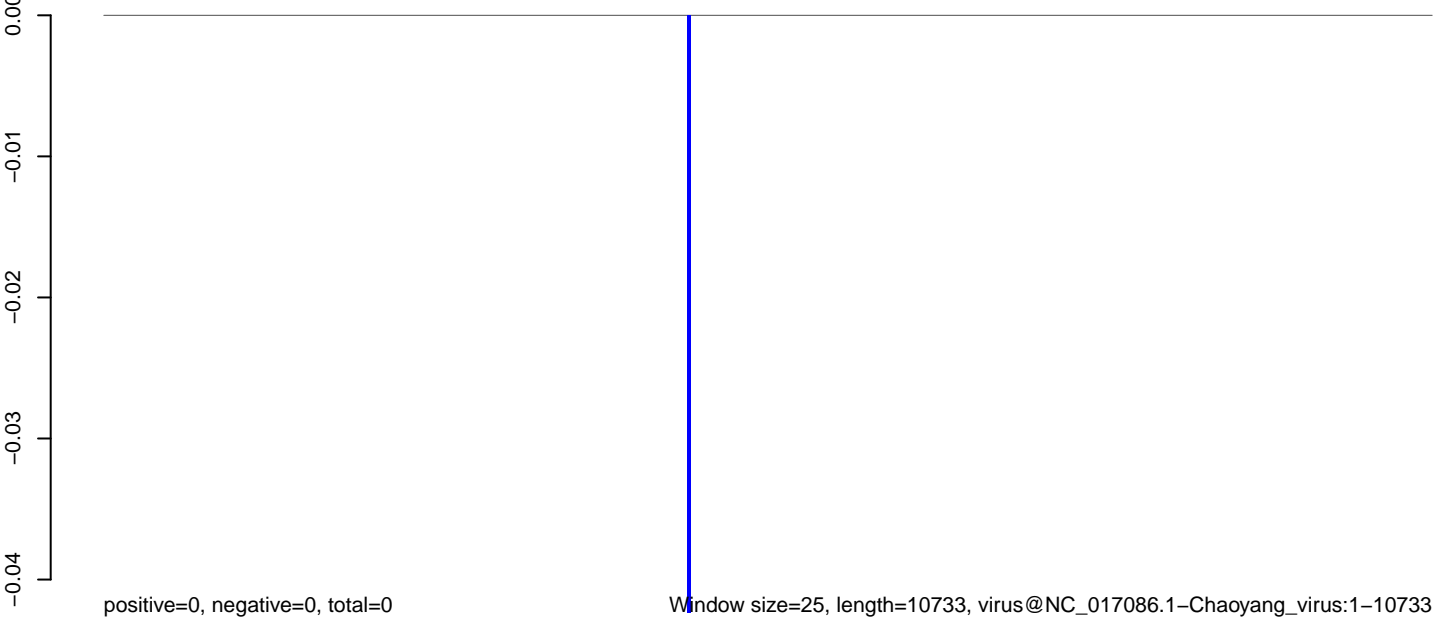
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



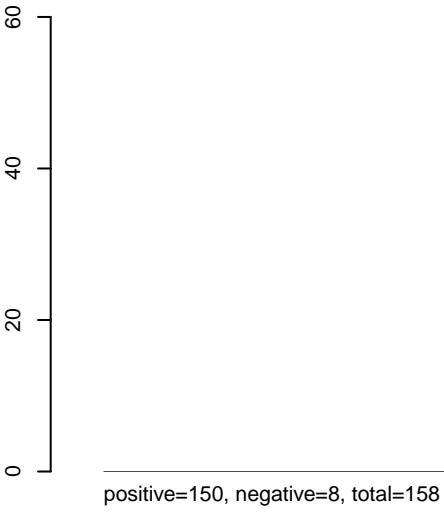
AeAlbo_C636_cells.rep



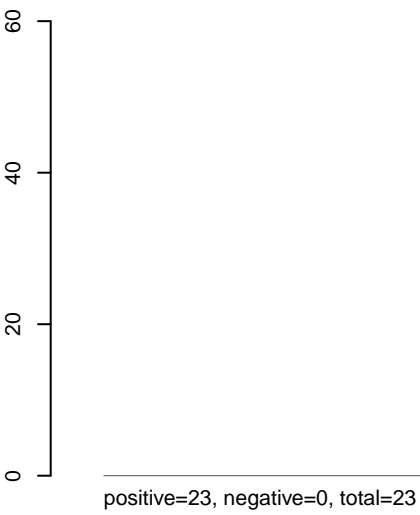
Window size=25, length=10733, virus@NC_017086.1-Chaoyang_virus:1-10733

0 2000 4000 6000 8000 10000

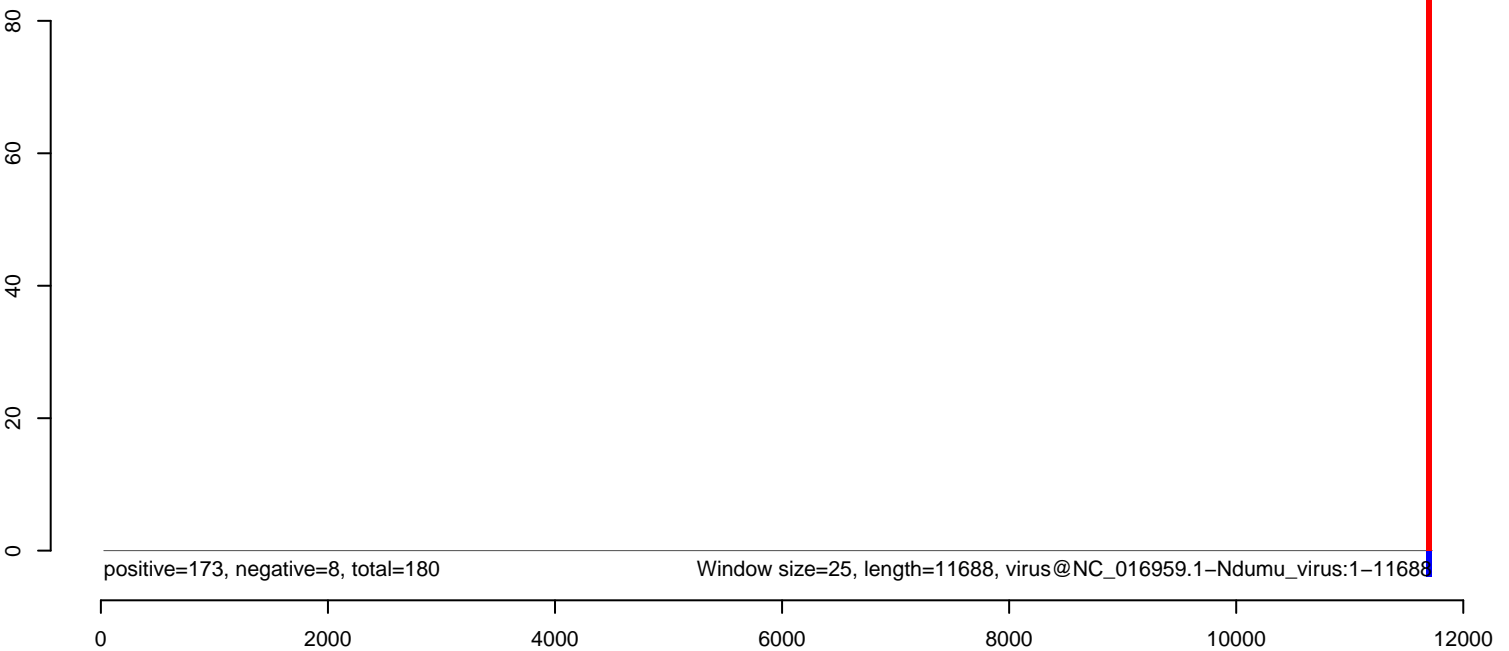
AeAlbo_C636_cells.18_23.rep



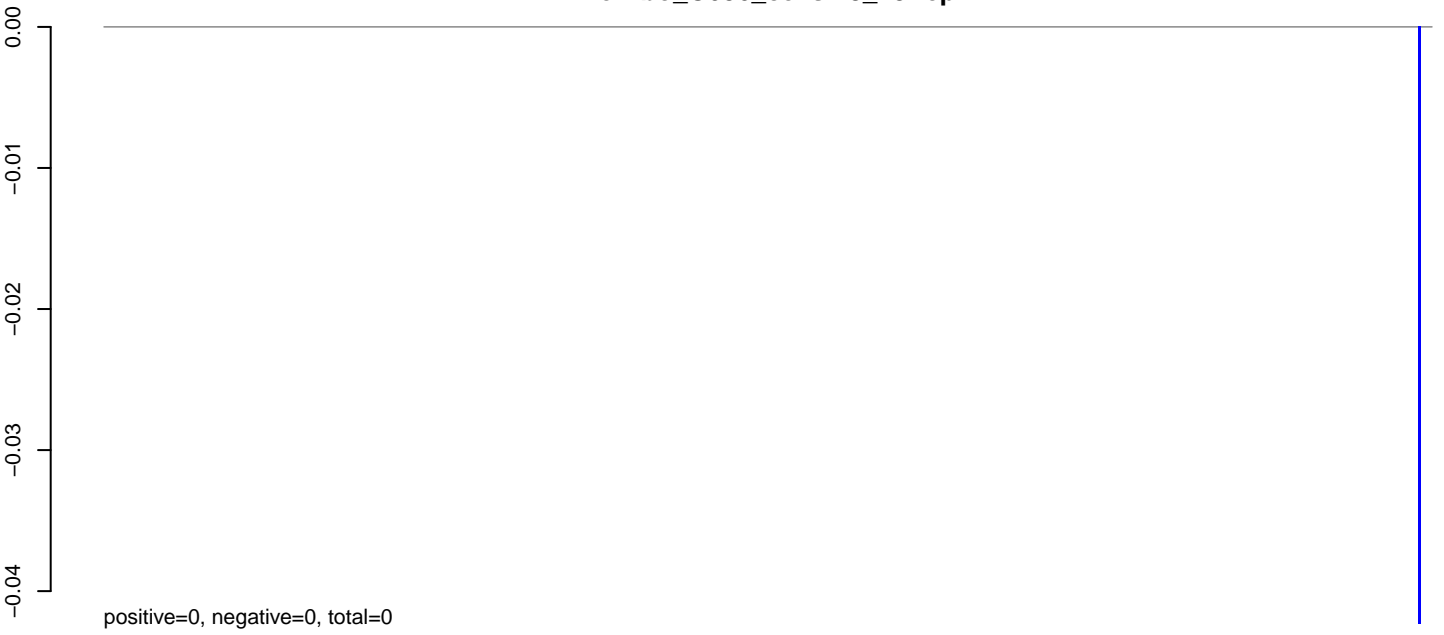
AeAlbo_C636_cells.24_35.rep



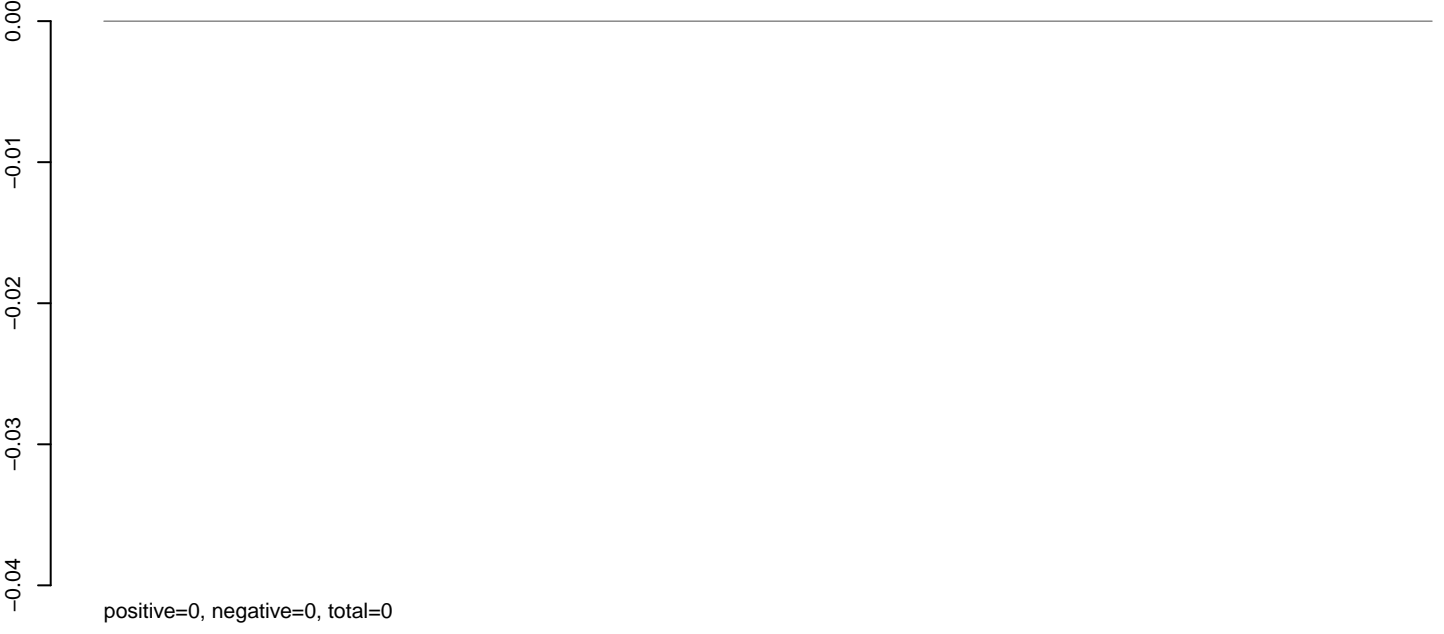
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



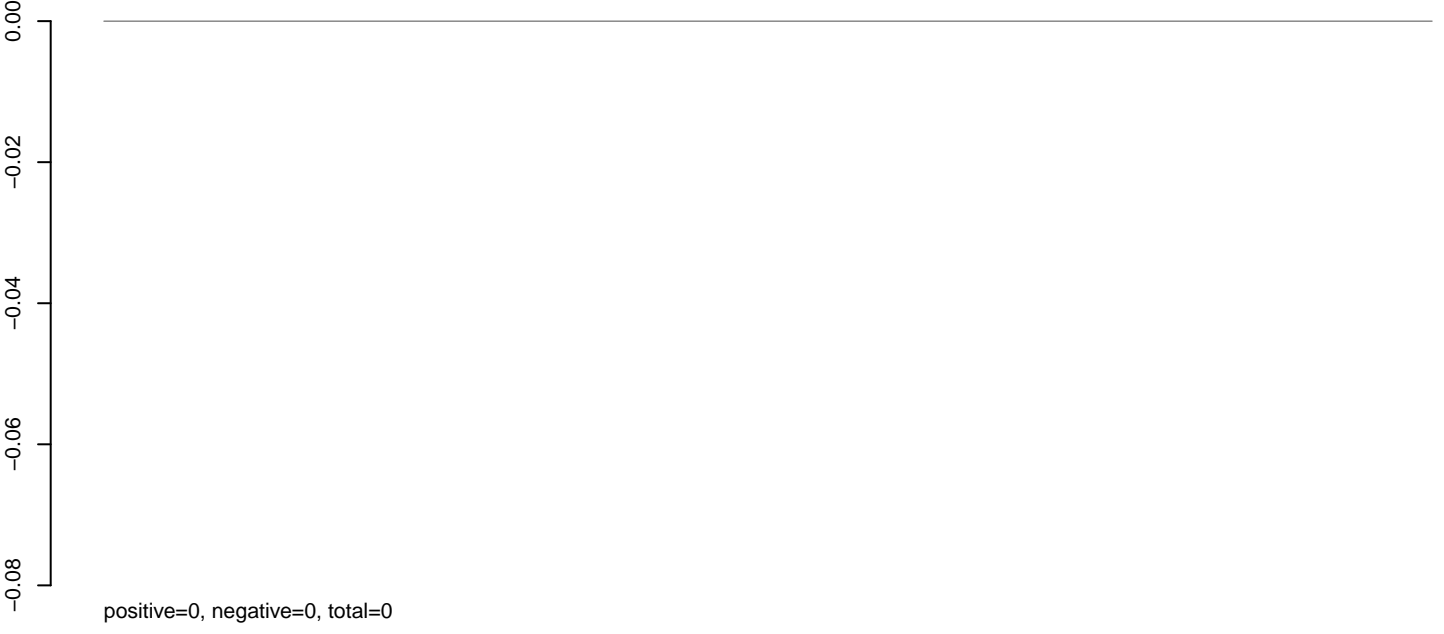
Window size=25, length=11381, virus@NC_013528.1-Fort_Morgan_virus:1-11381

0 2000 4000 6000 8000 10000

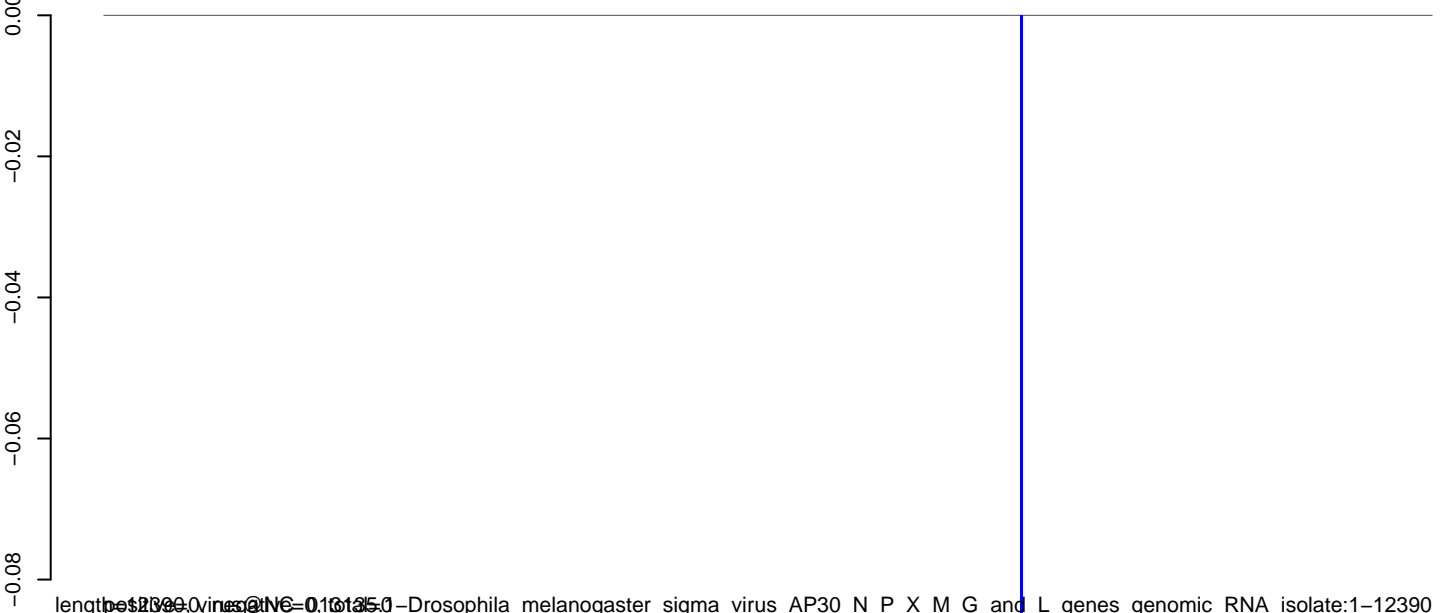
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



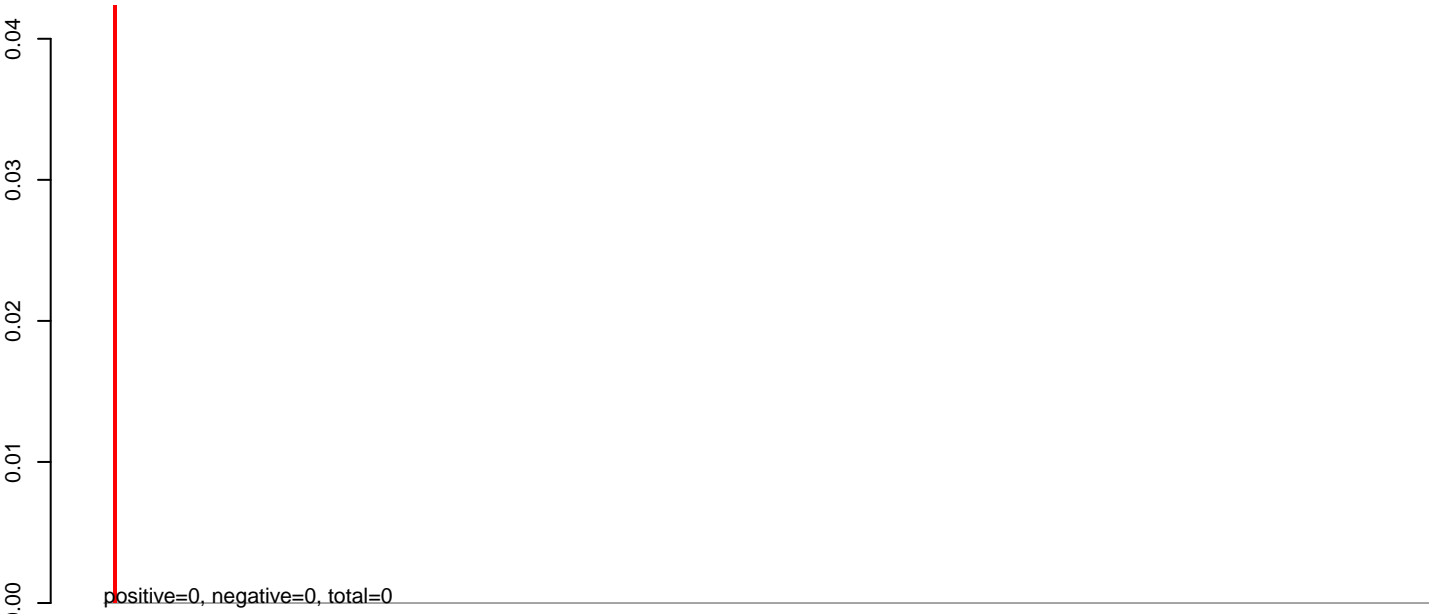
AeAlbo_C636_cells.rep



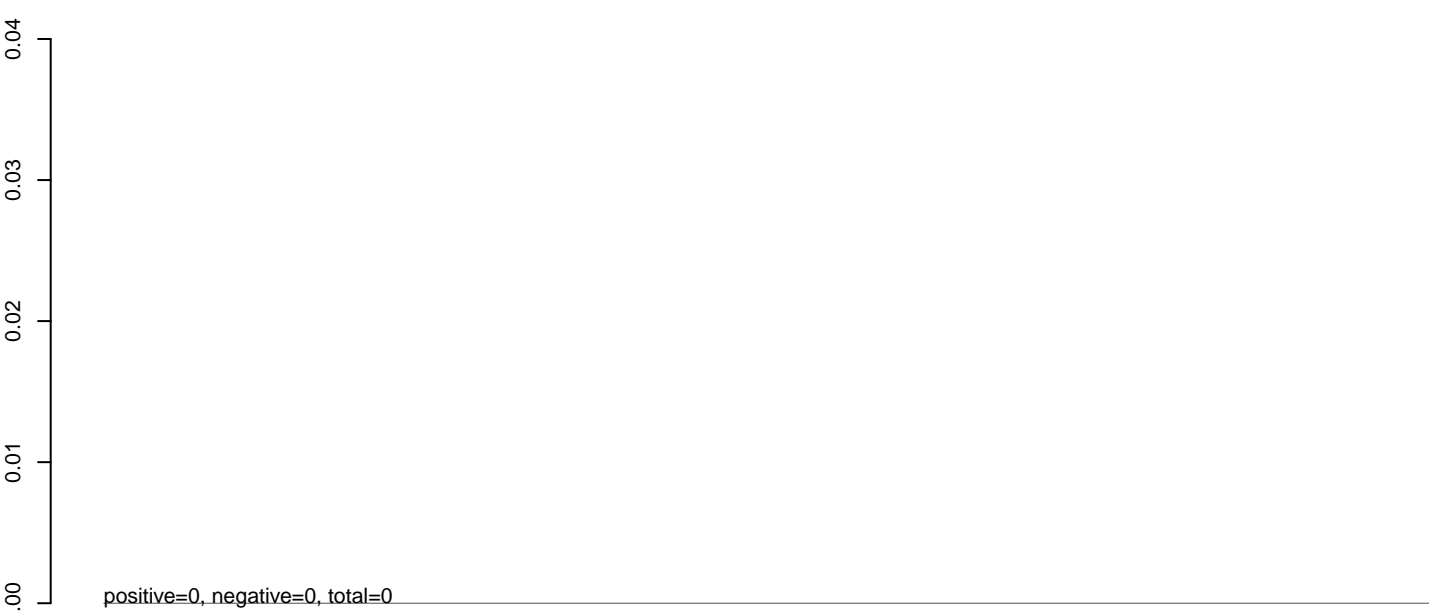
lengths: 2399, 16135, 16135 -Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390

0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

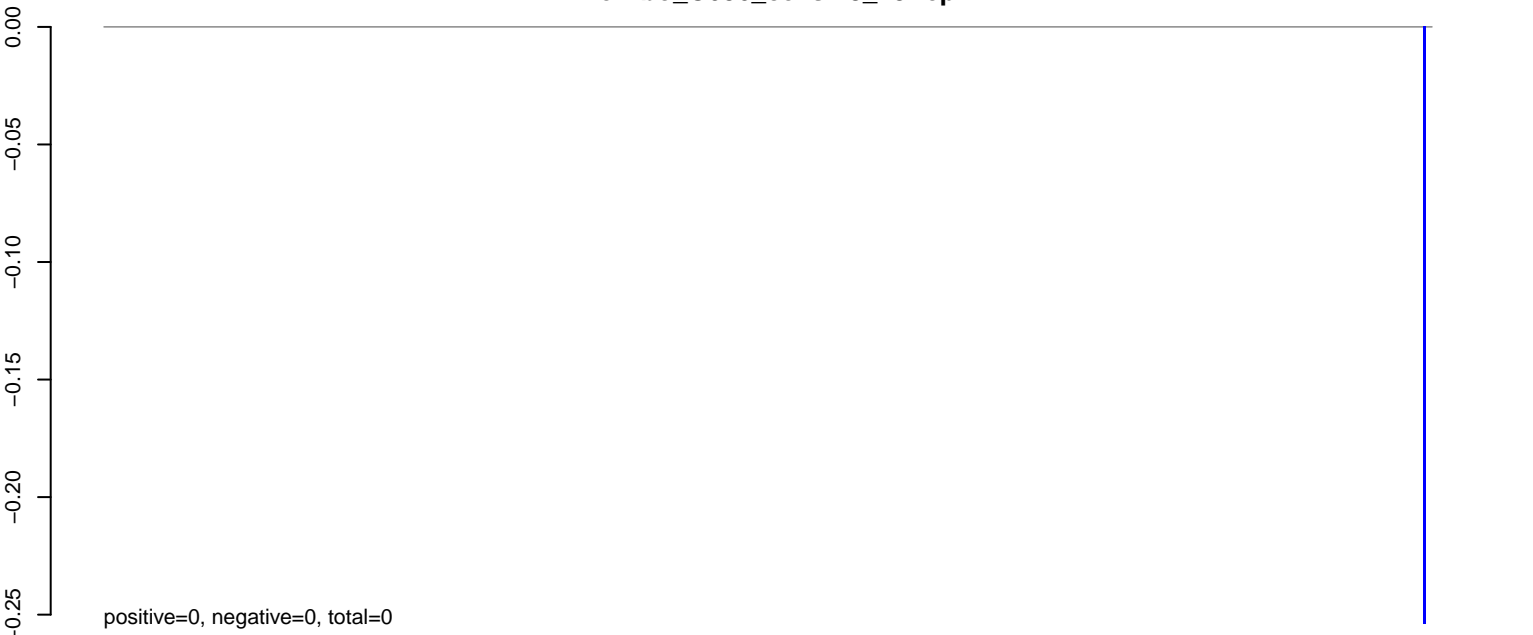


AeAlbo_C636_cells.rep

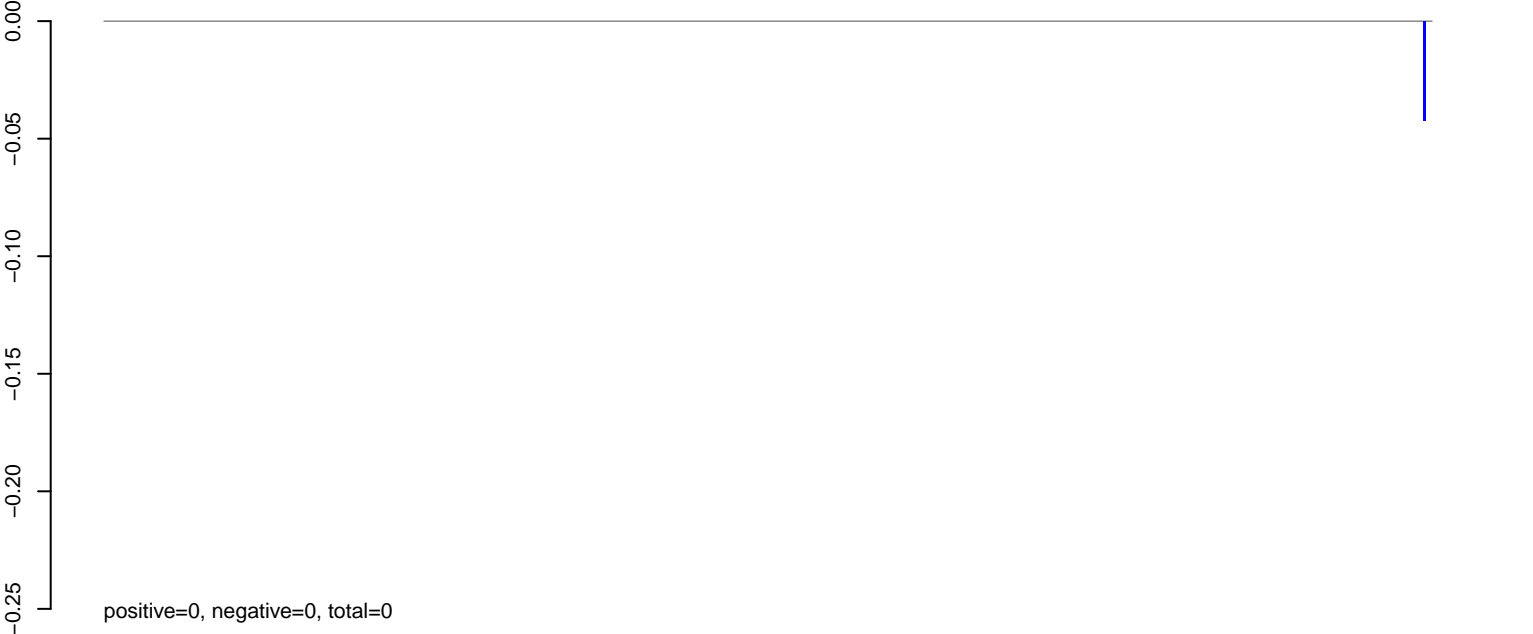


0 2000 4000 6000 8000 10000

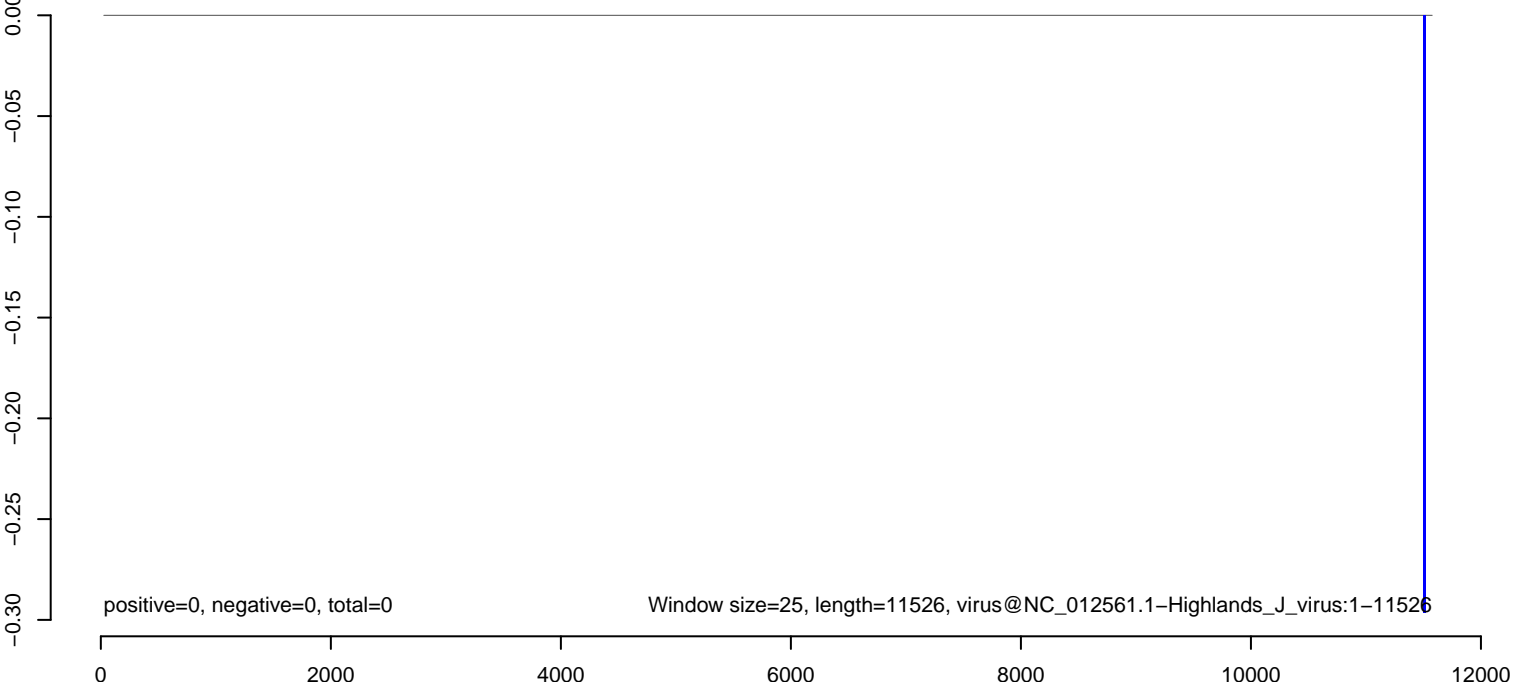
AeAlbo_C636_cells.18_23.rep



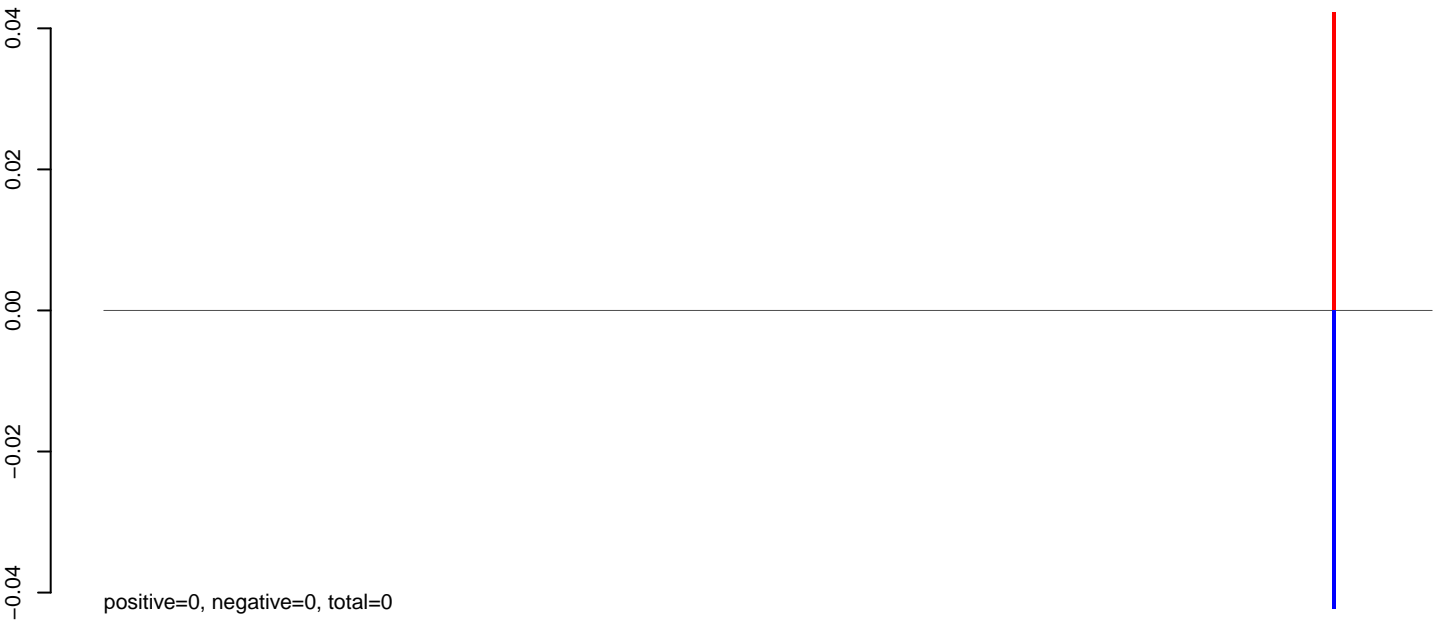
AeAlbo_C636_cells.24_35.rep



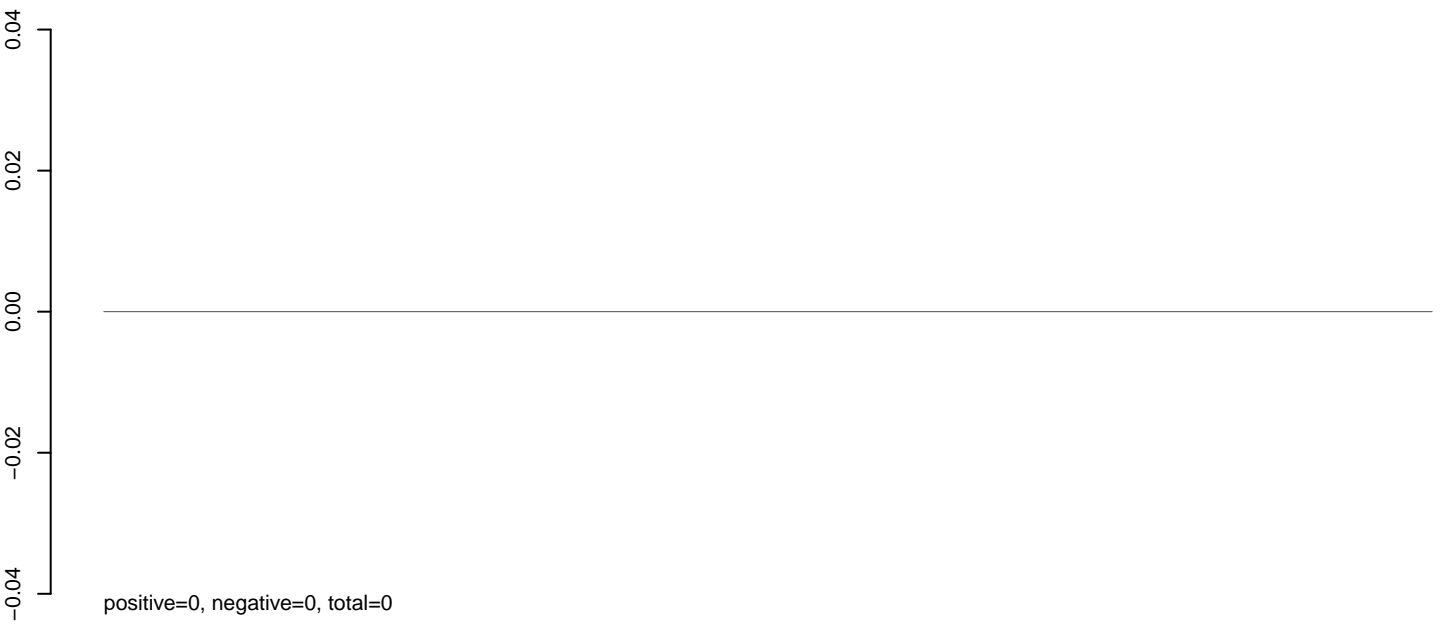
AeAlbo_C636_cells.rep



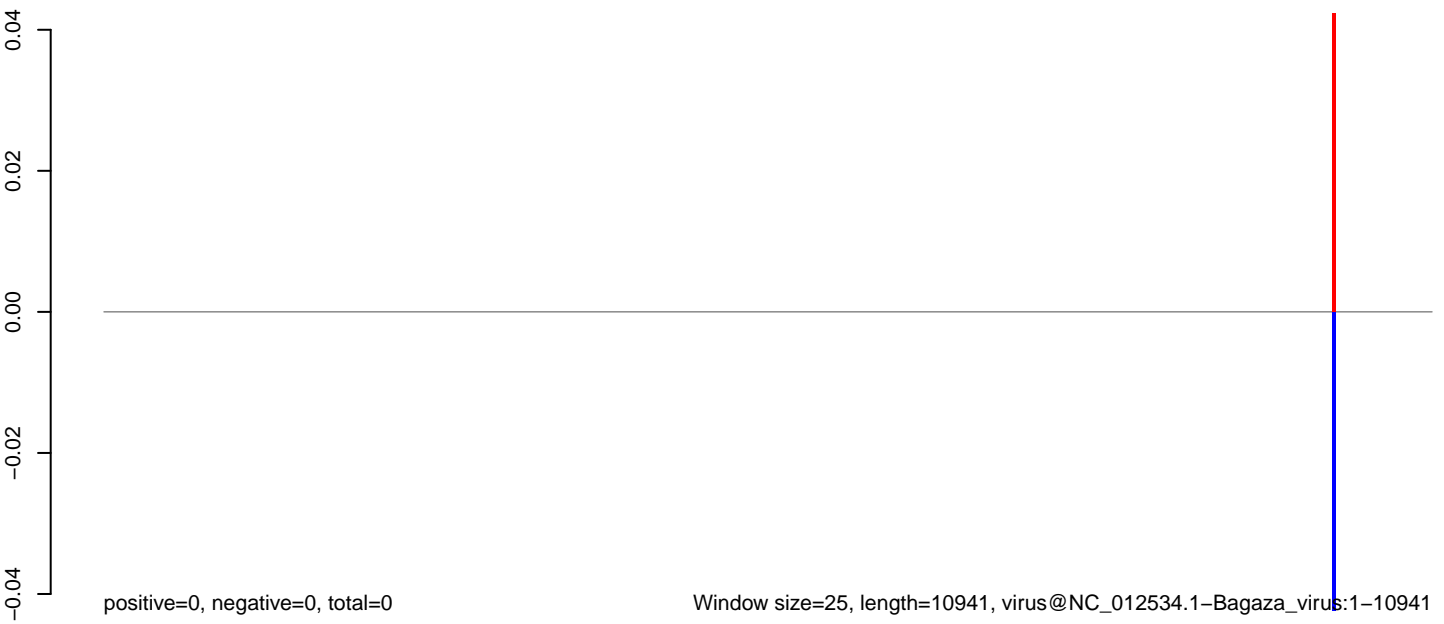
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

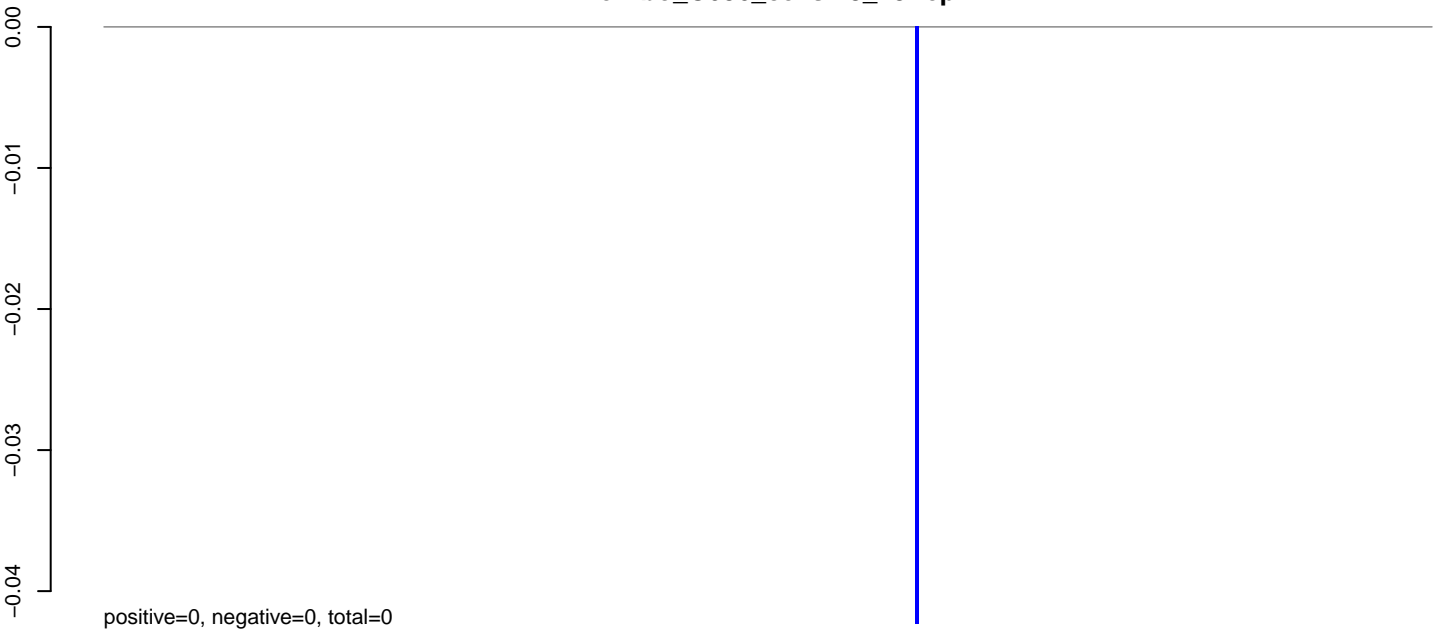


AeAlbo_C636_cells.rep

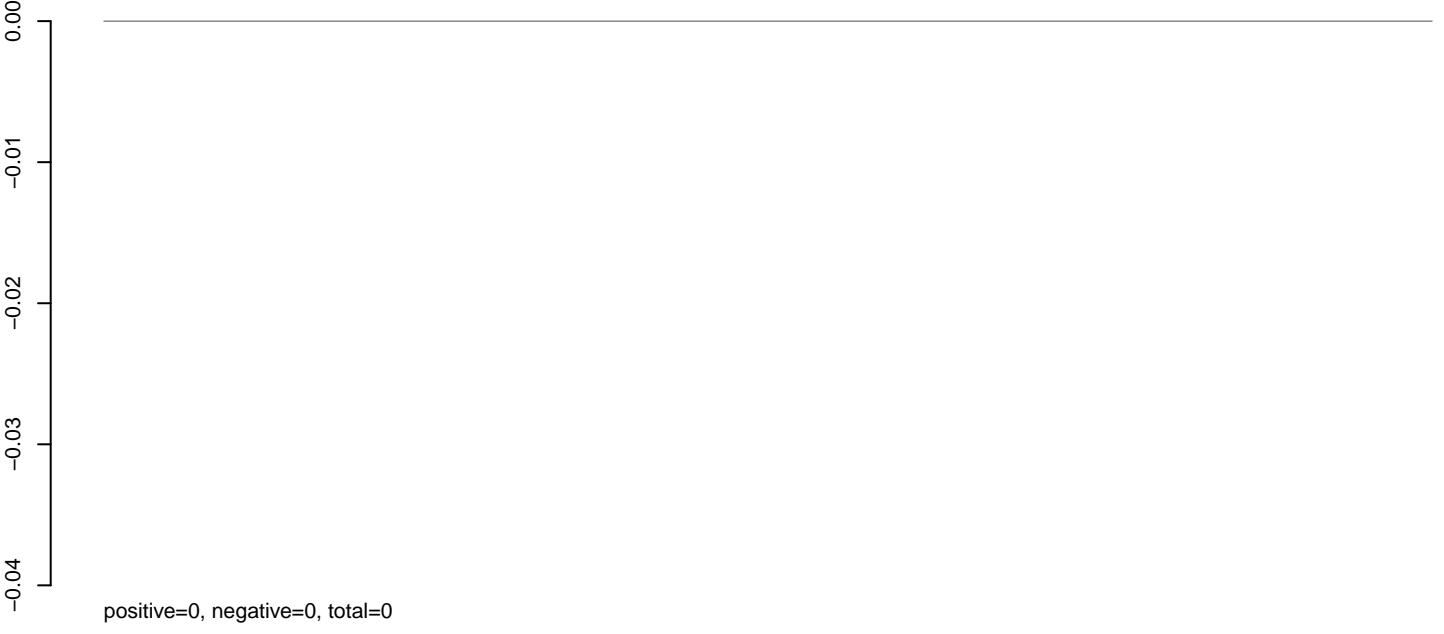


0 2000 4000 6000 8000 10000

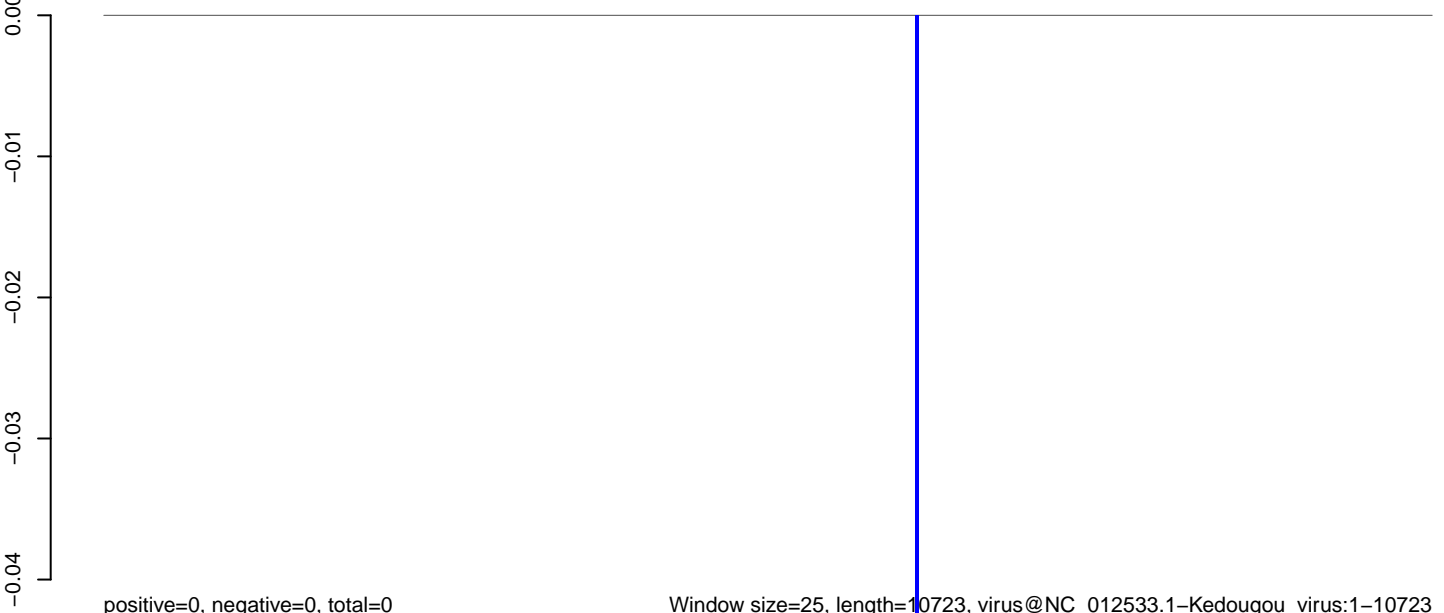
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



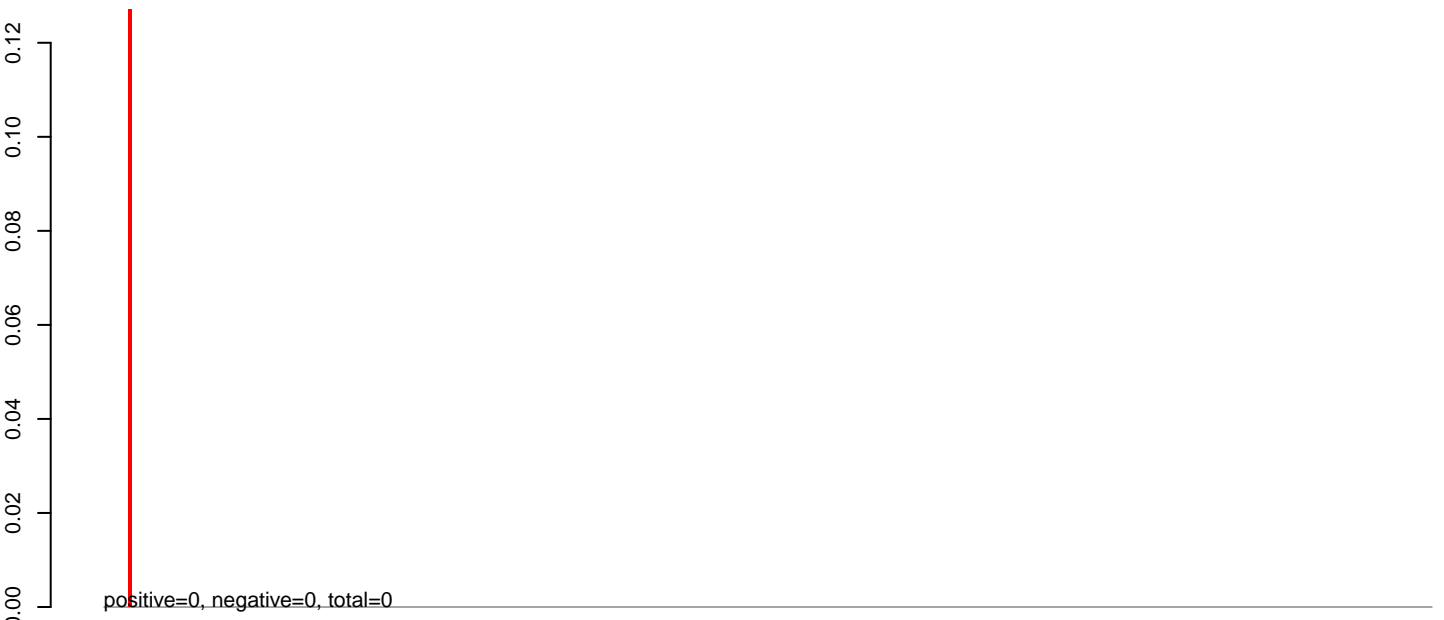
AeAlbo_C636_cells.rep



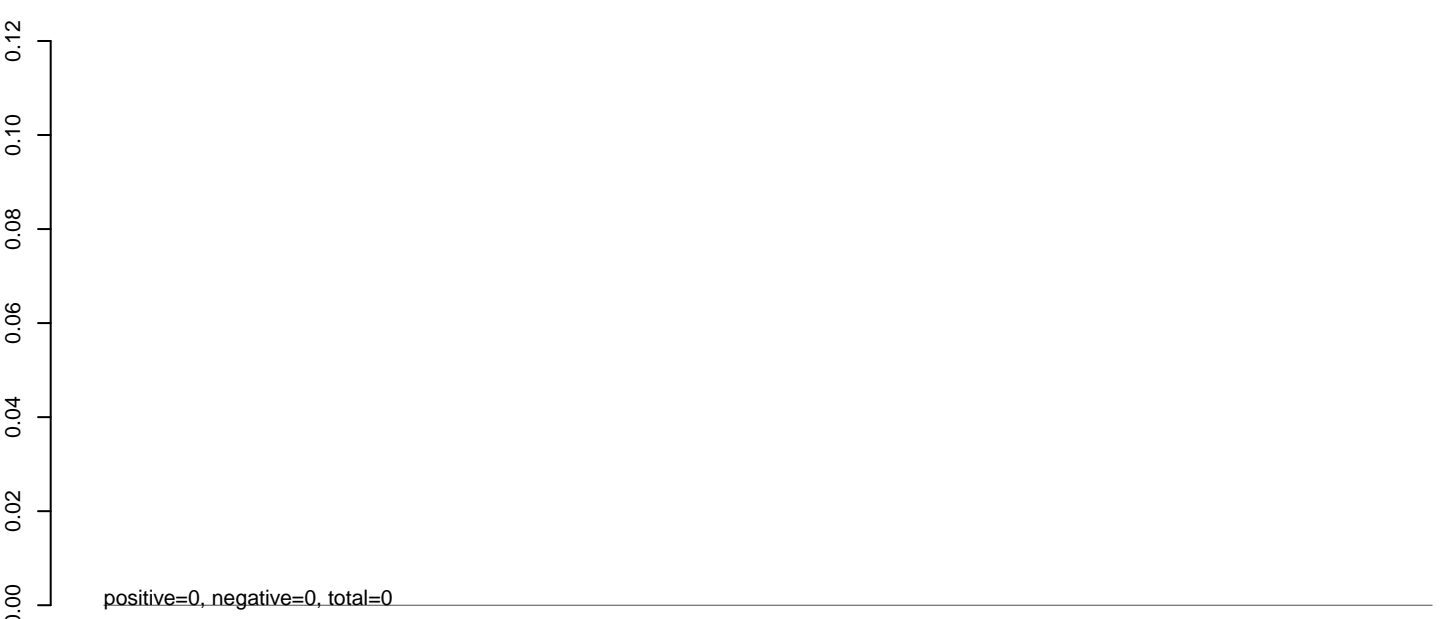
Window size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

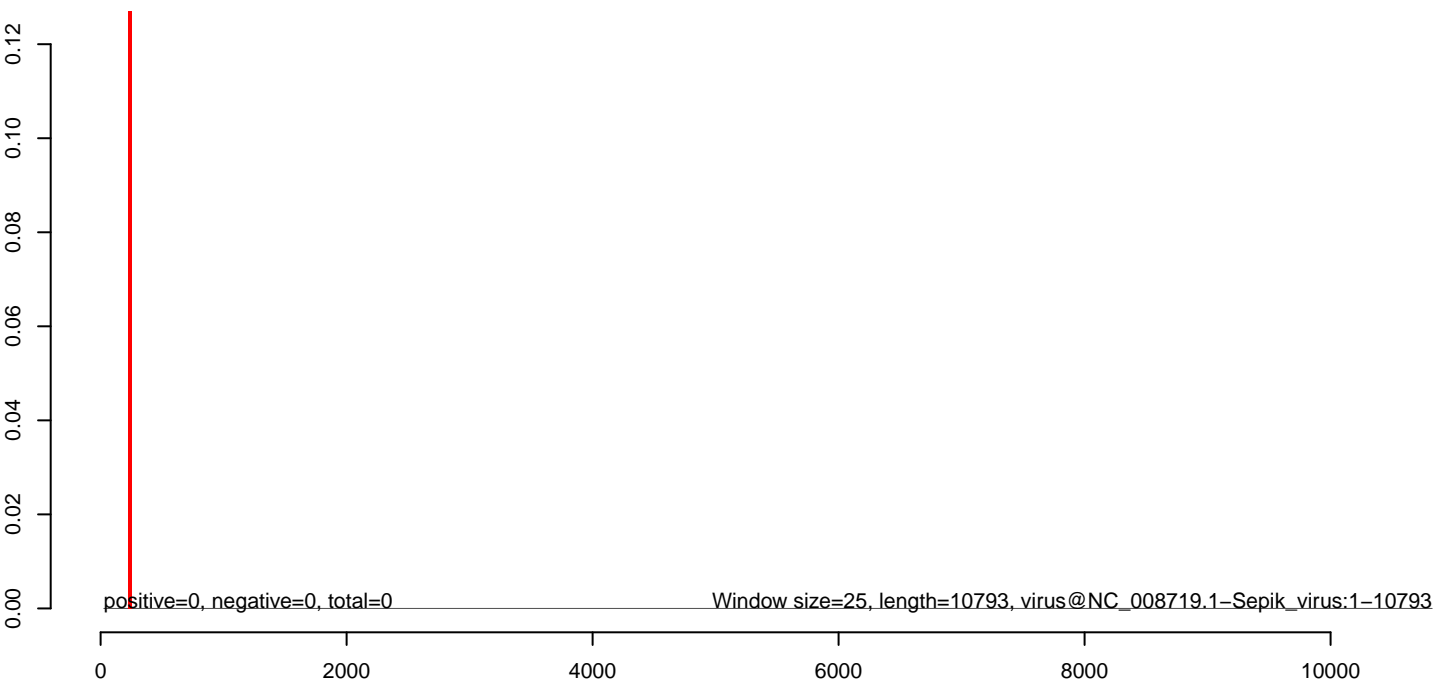
AeAlbo_C636_cells.18_23.rep



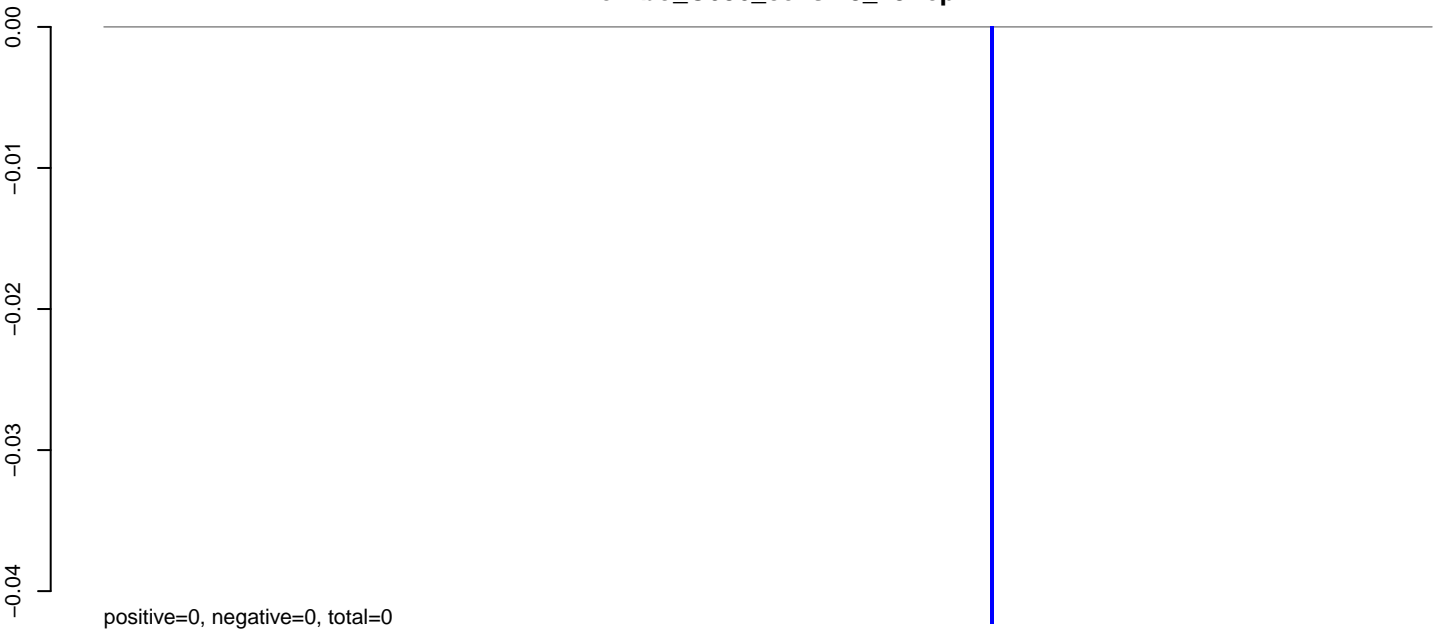
AeAlbo_C636_cells.24_35.rep



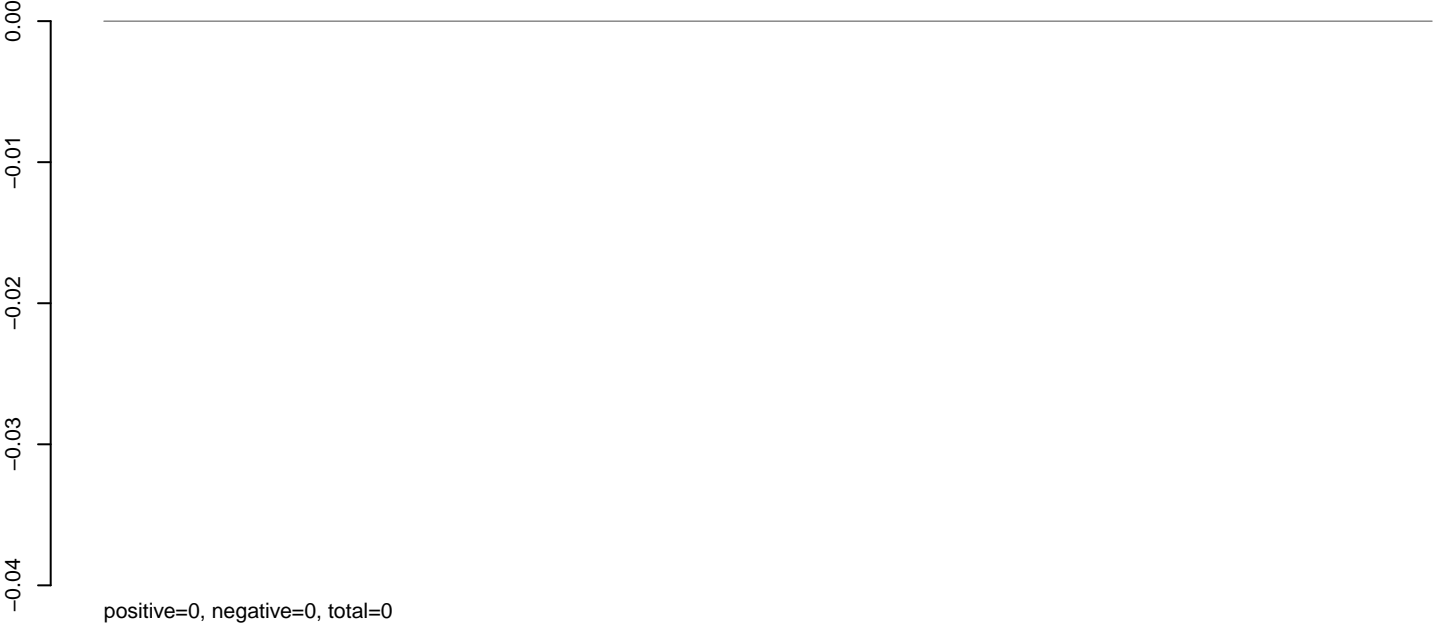
AeAlbo_C636_cells.rep



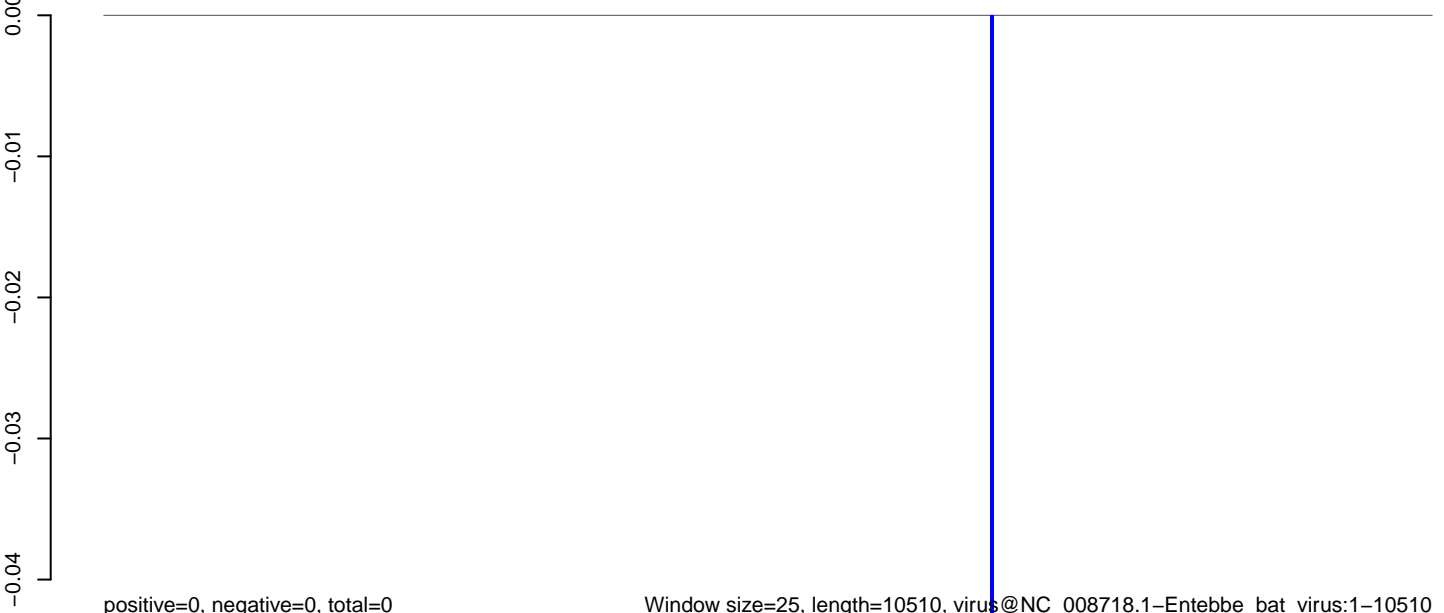
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



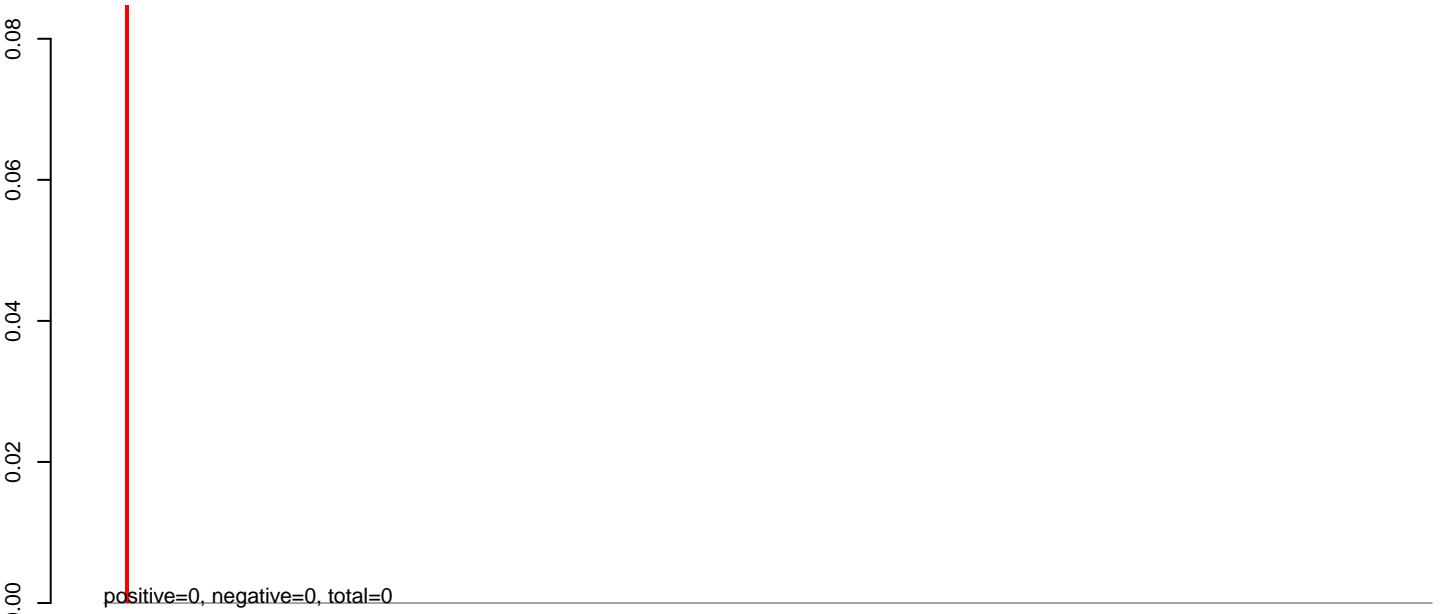
AeAlbo_C636_cells.rep



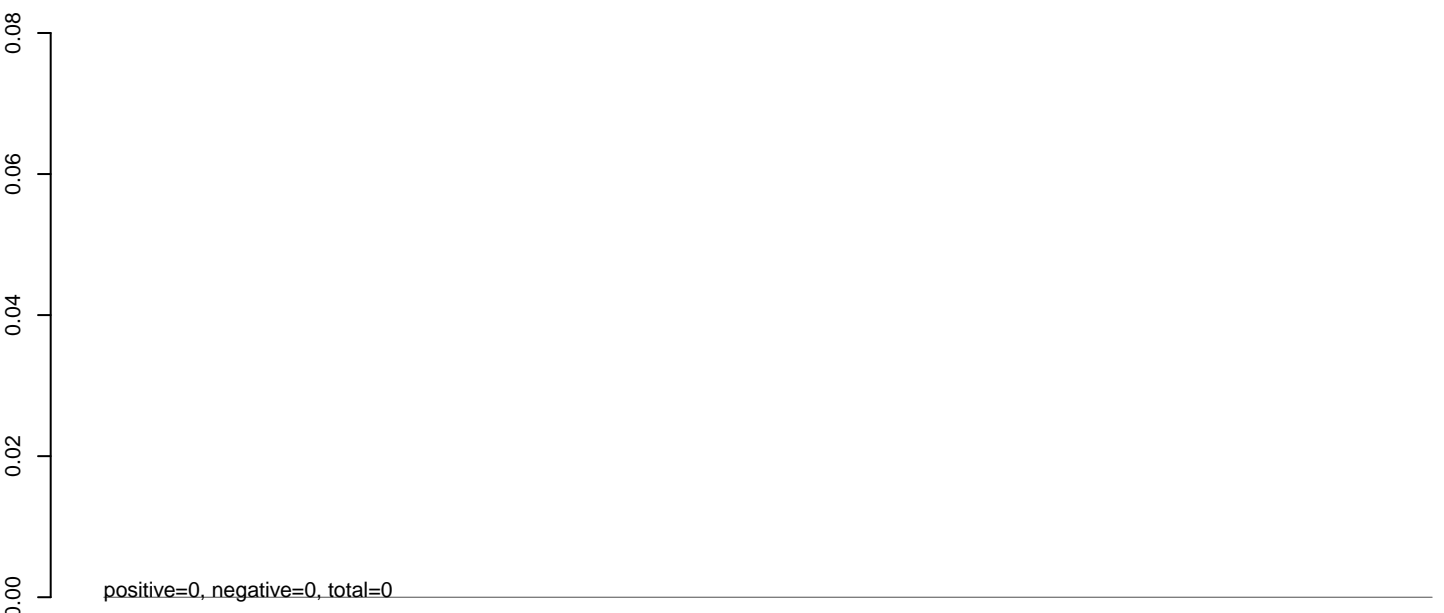
Window size=25, length=10510, virus@NC_008718.1-Entebbe_bat_virus:1-10510

0 2000 4000 6000 8000 10000

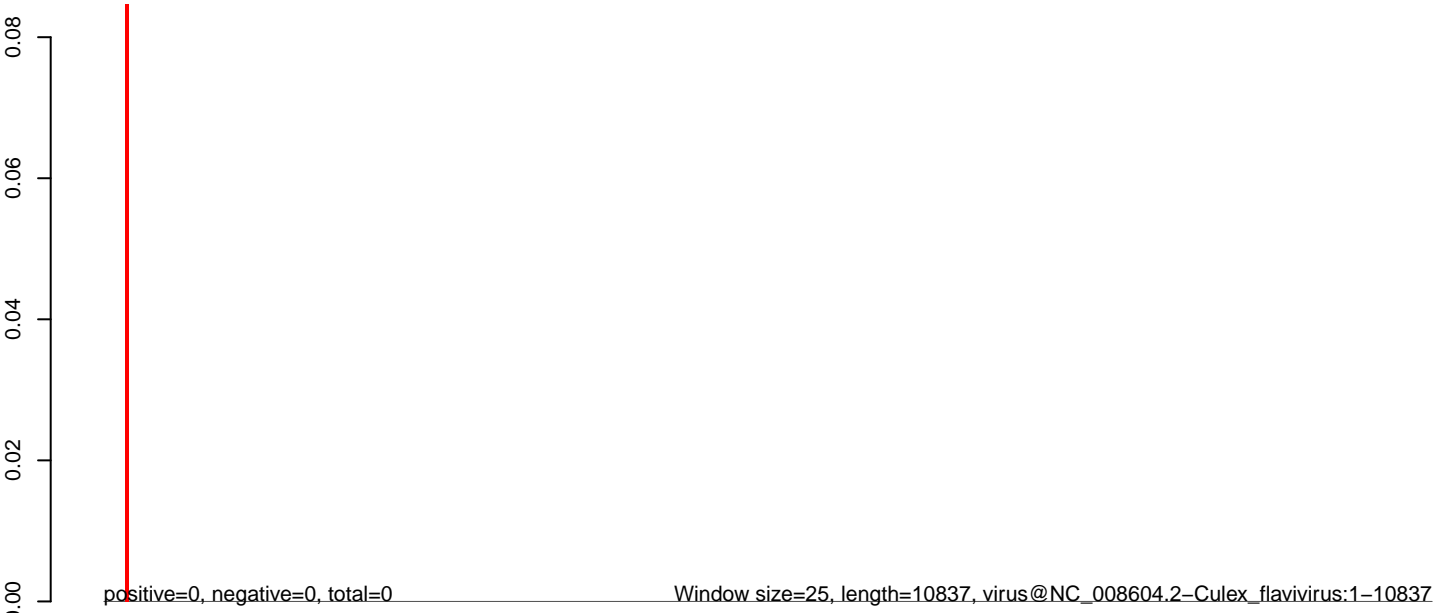
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

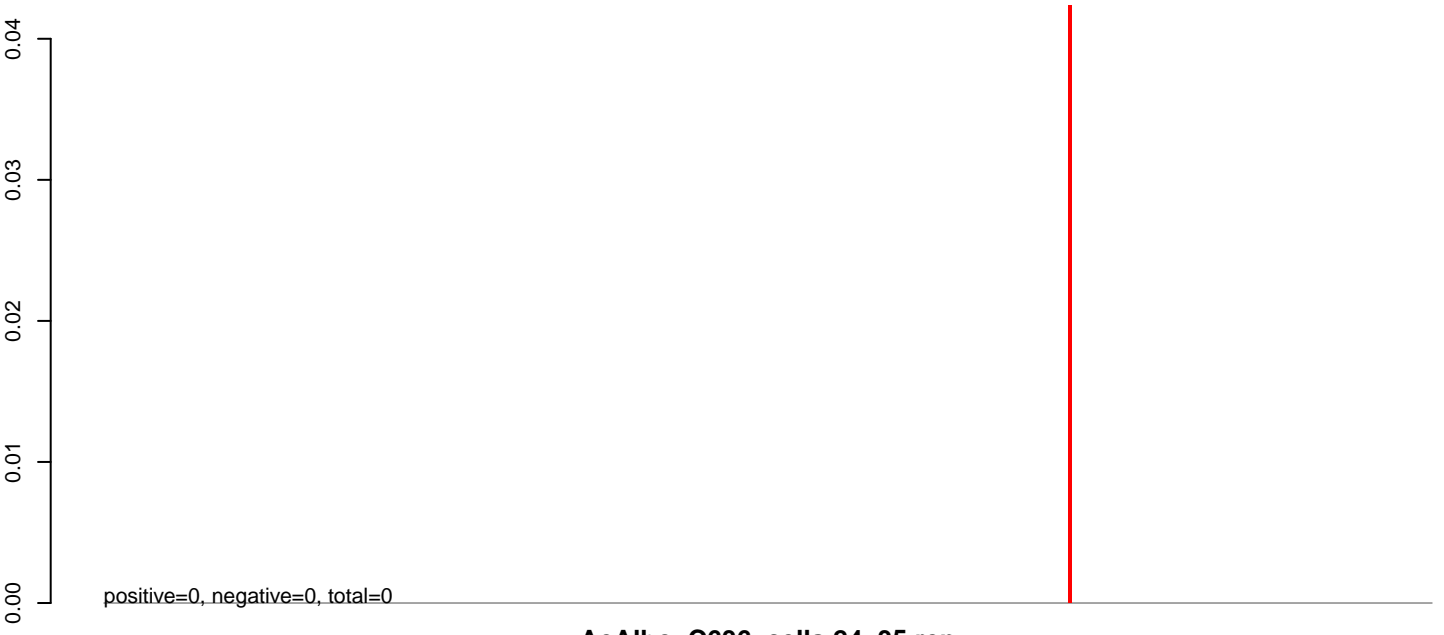


AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

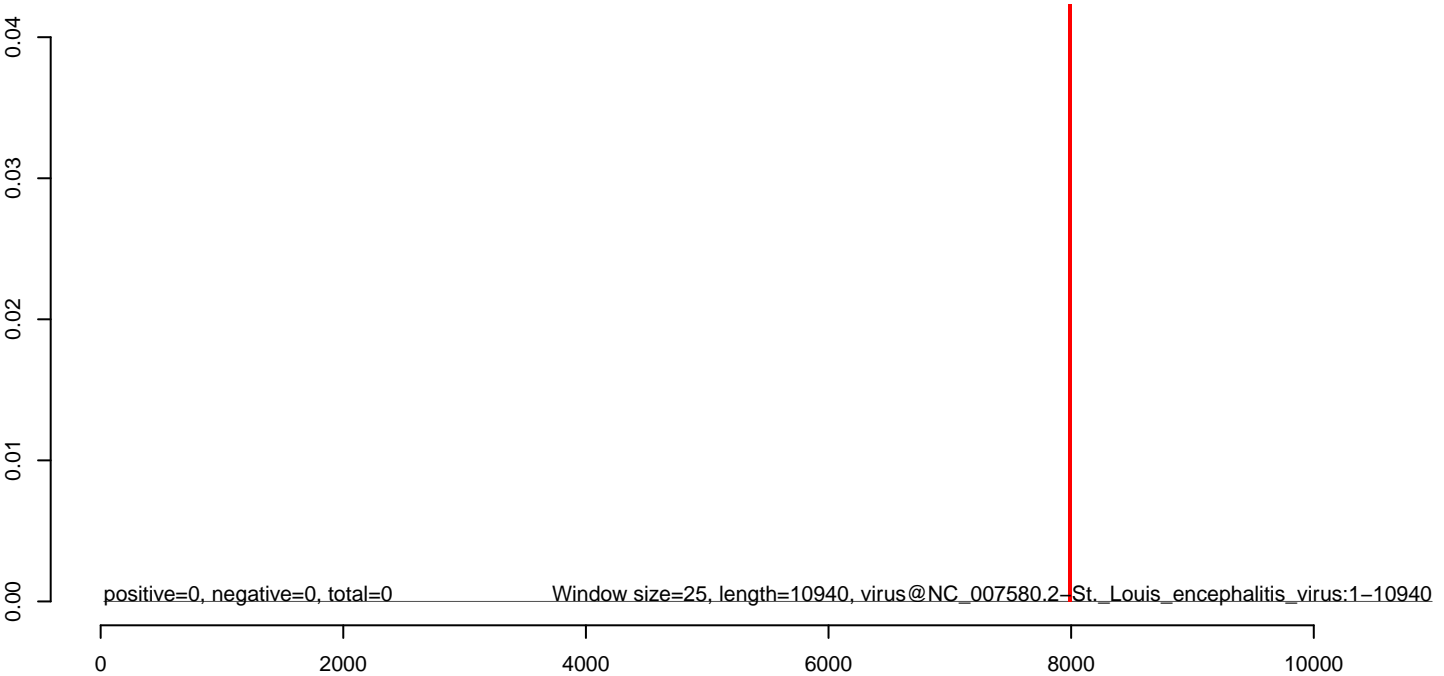
AeAlbo_C636_cells.18_23.rep



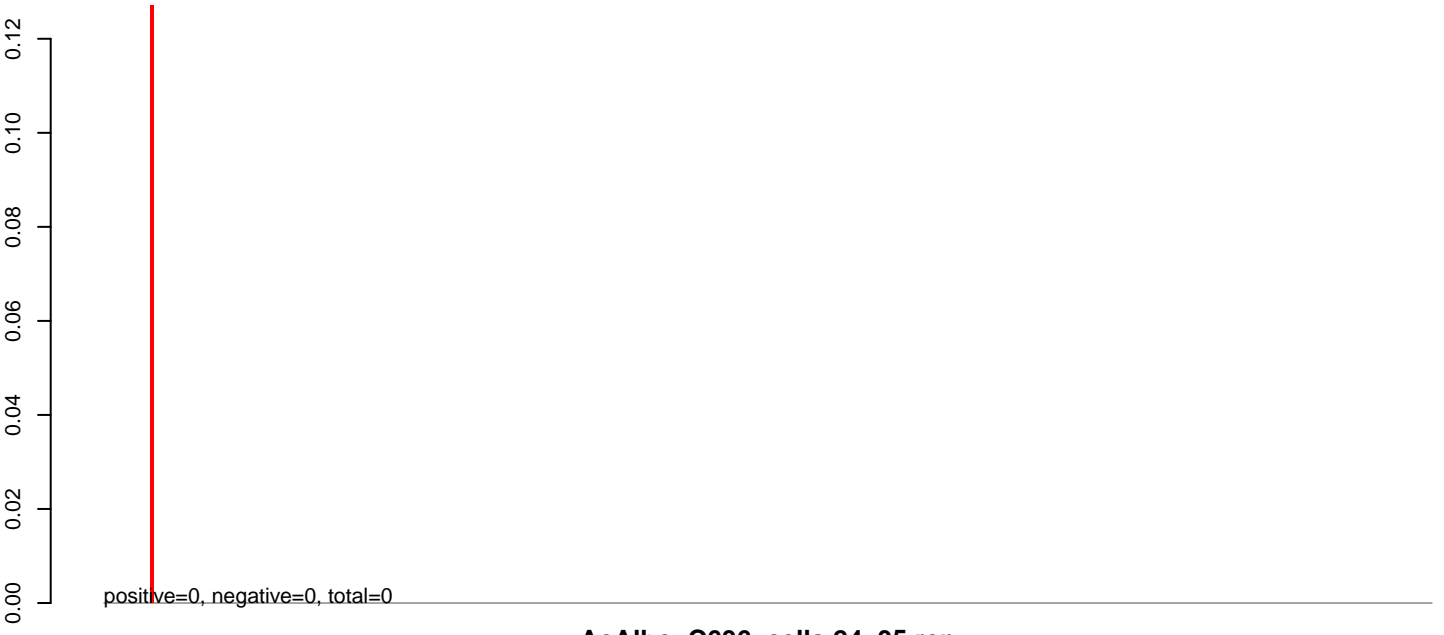
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



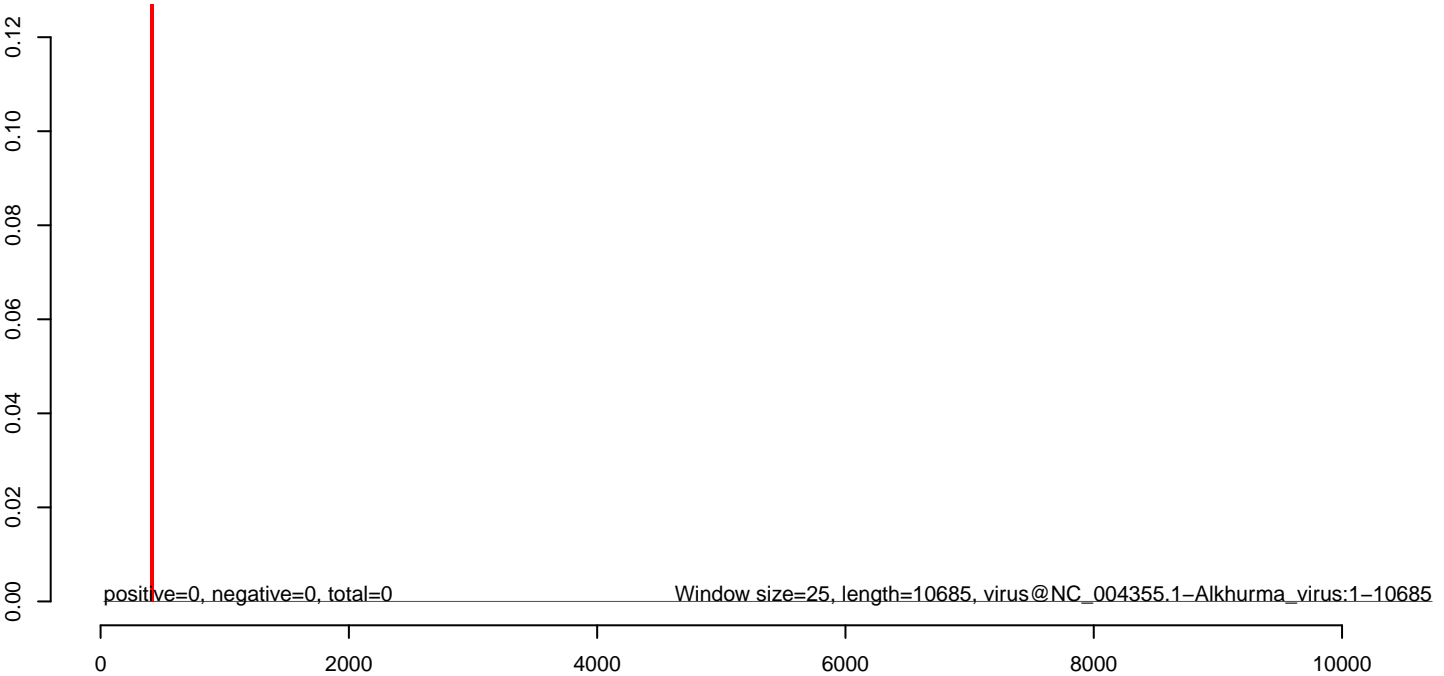
AeAlbo_C636_cells.18_23.rep



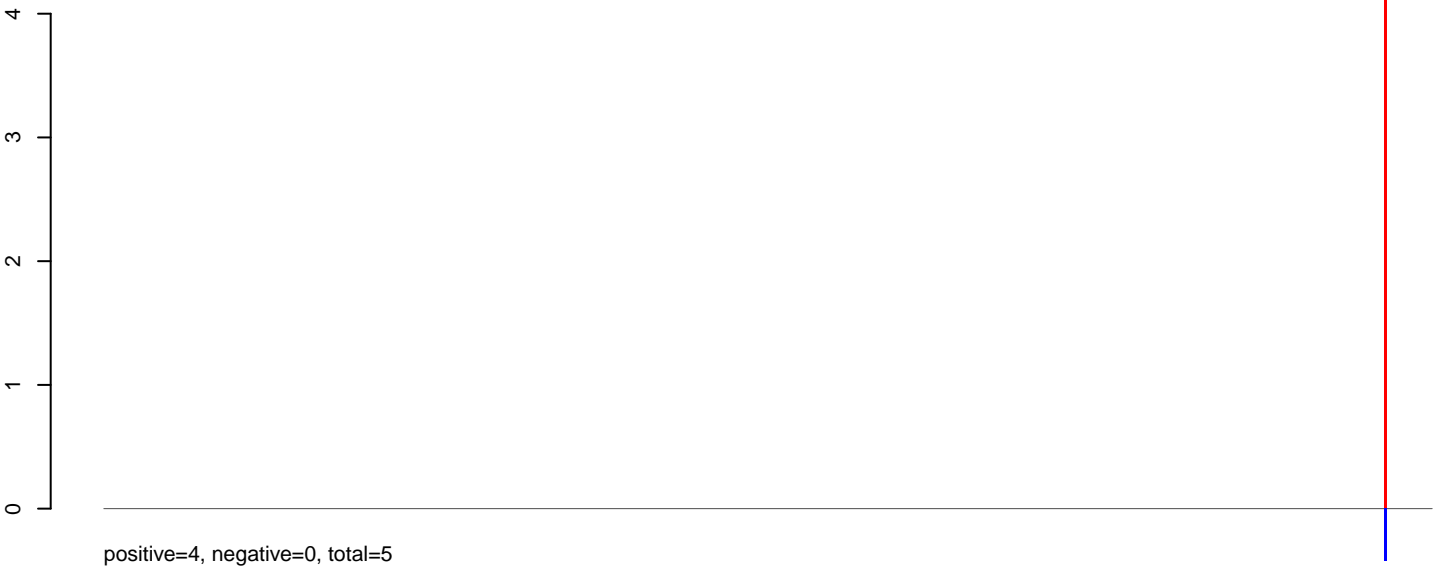
AeAlbo_C636_cells.24_35.rep



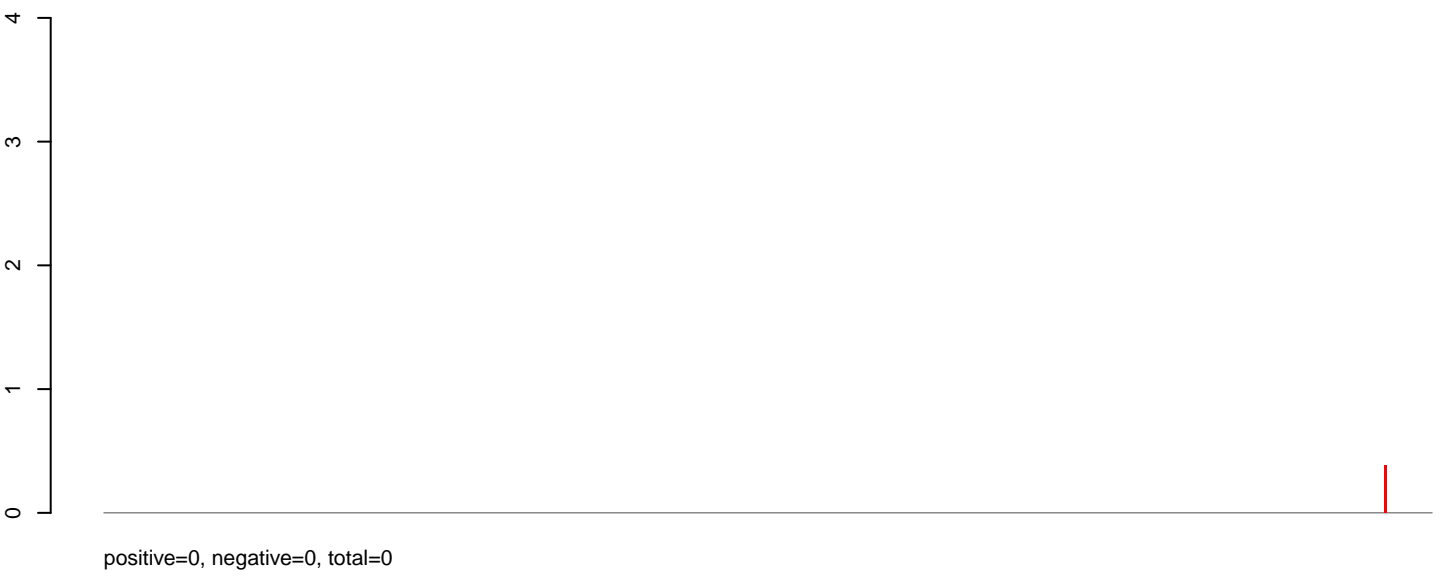
AeAlbo_C636_cells.rep



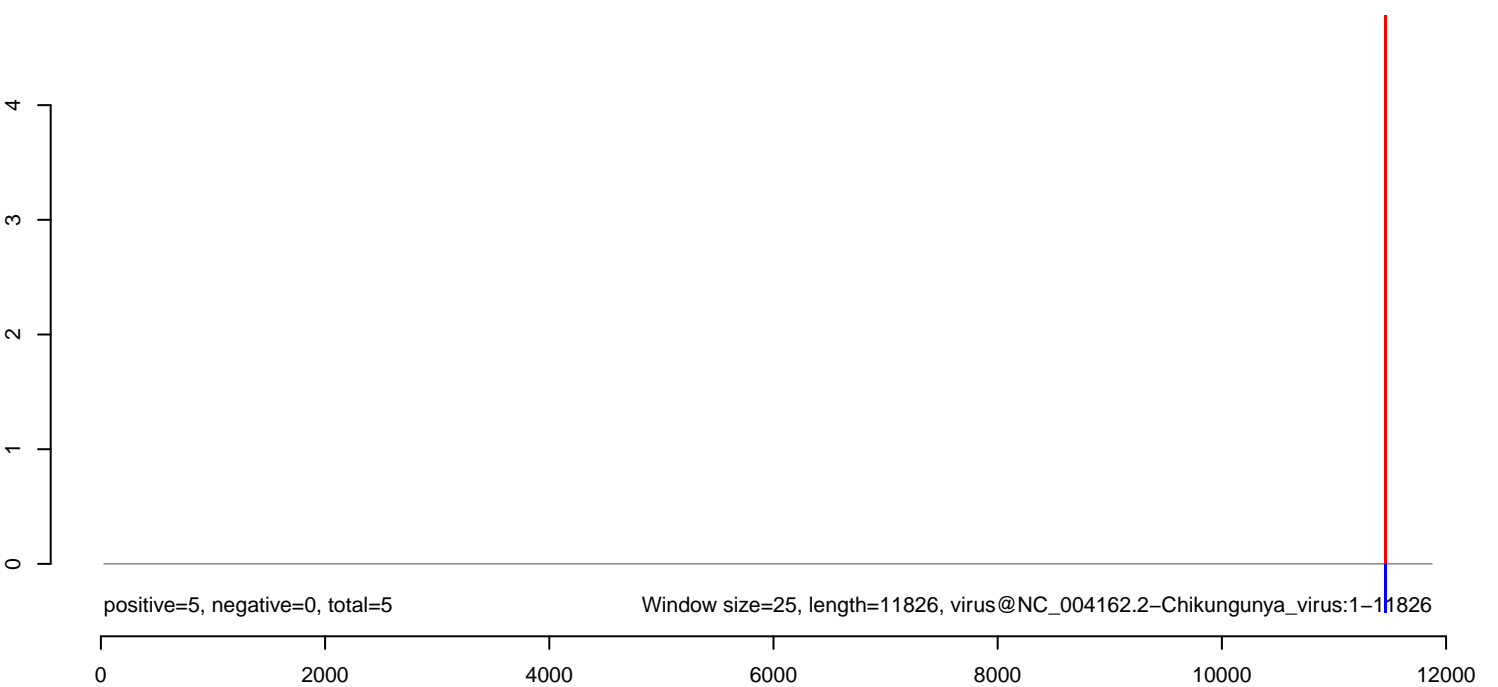
AeAlbo_C636_cells.18_23.rep



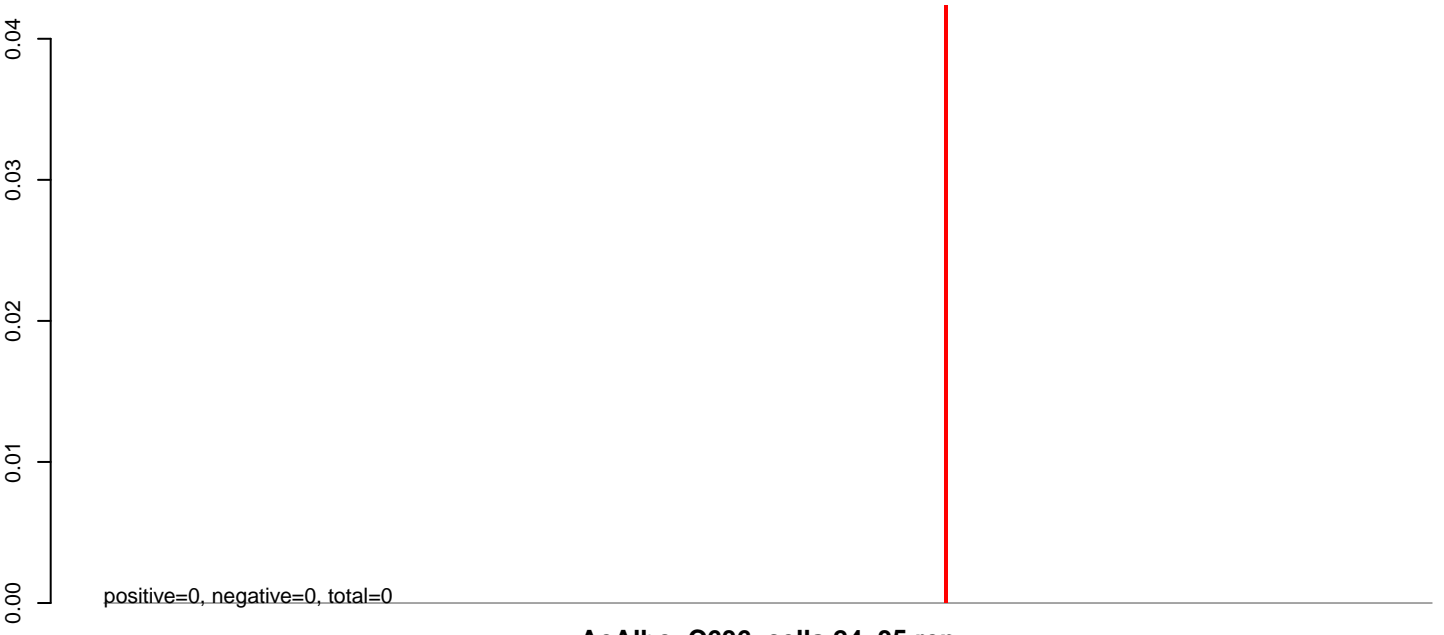
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



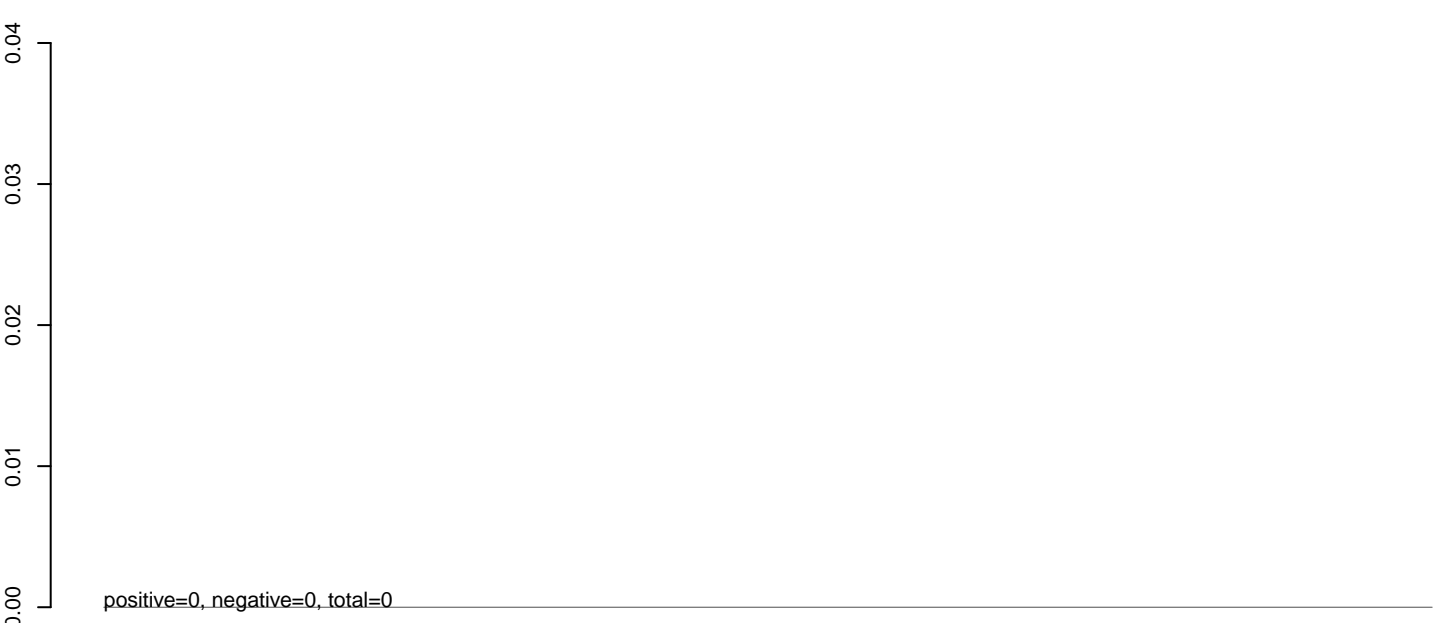
AeAlbo_C636_cells.rep



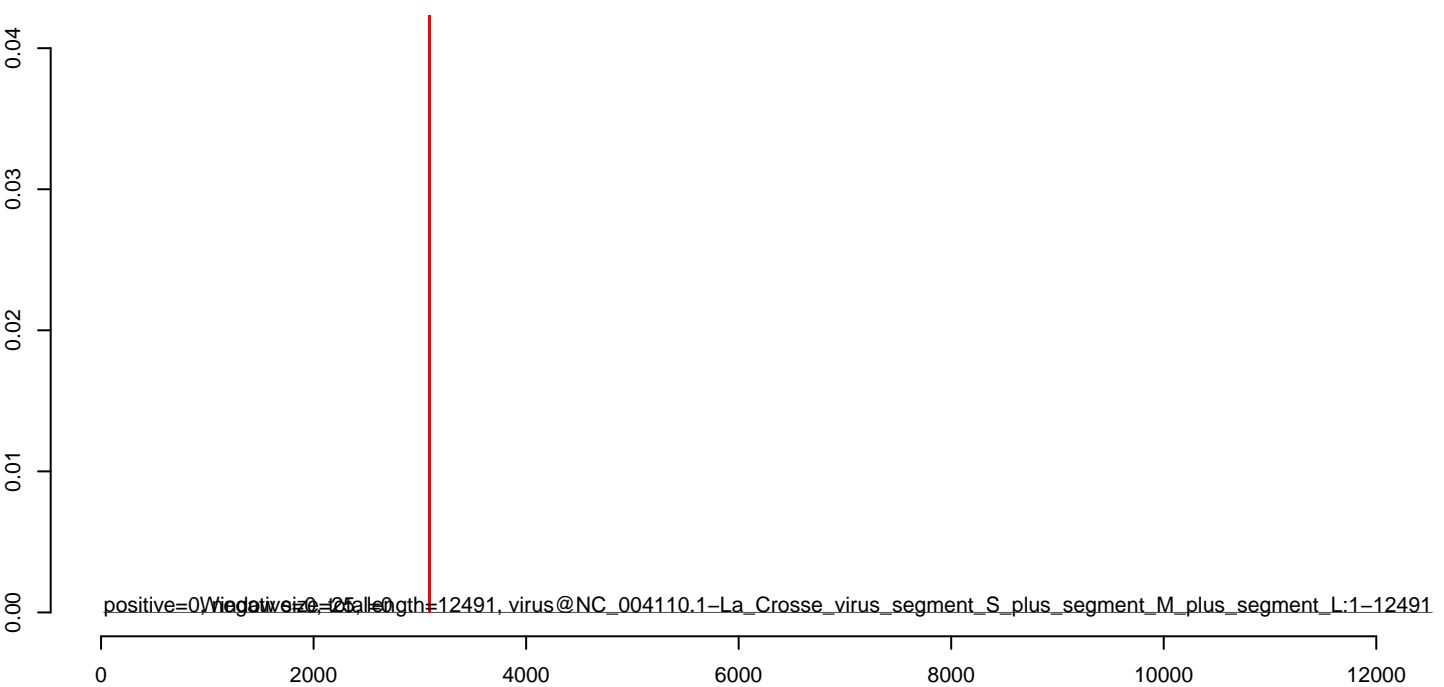
AeAlbo_C636_cells.18_23.rep



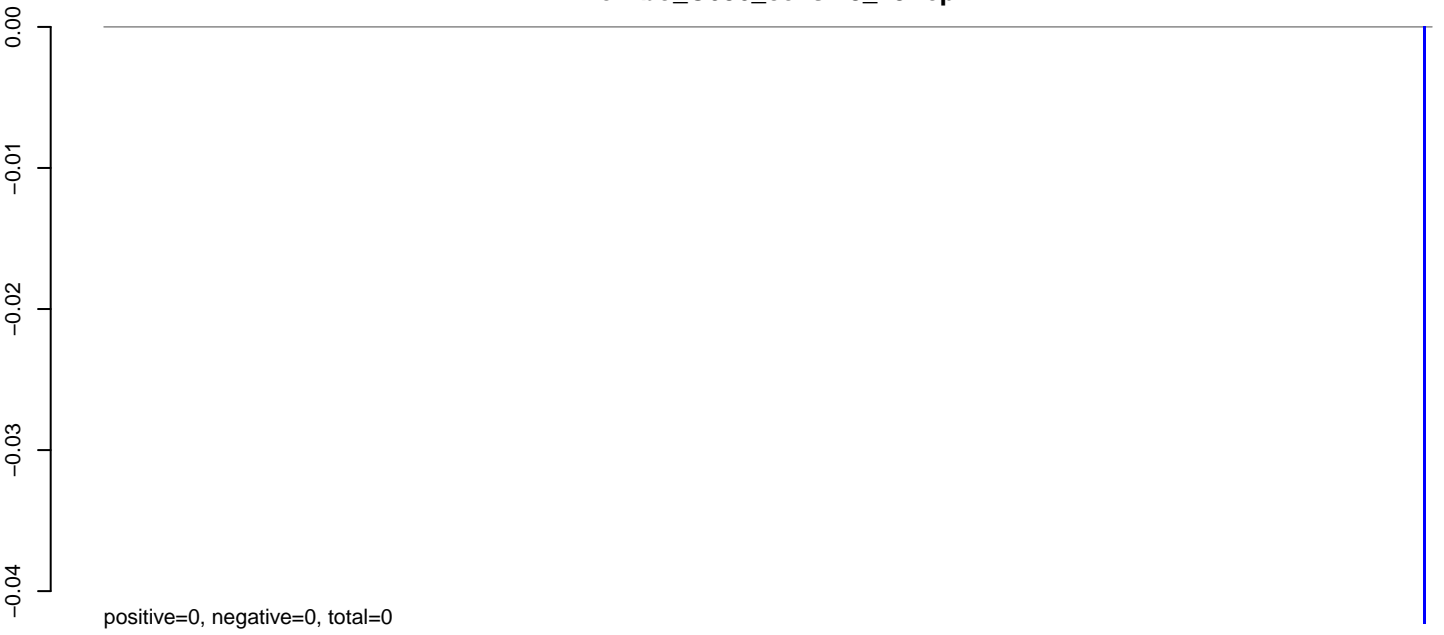
AeAlbo_C636_cells.24_35.rep



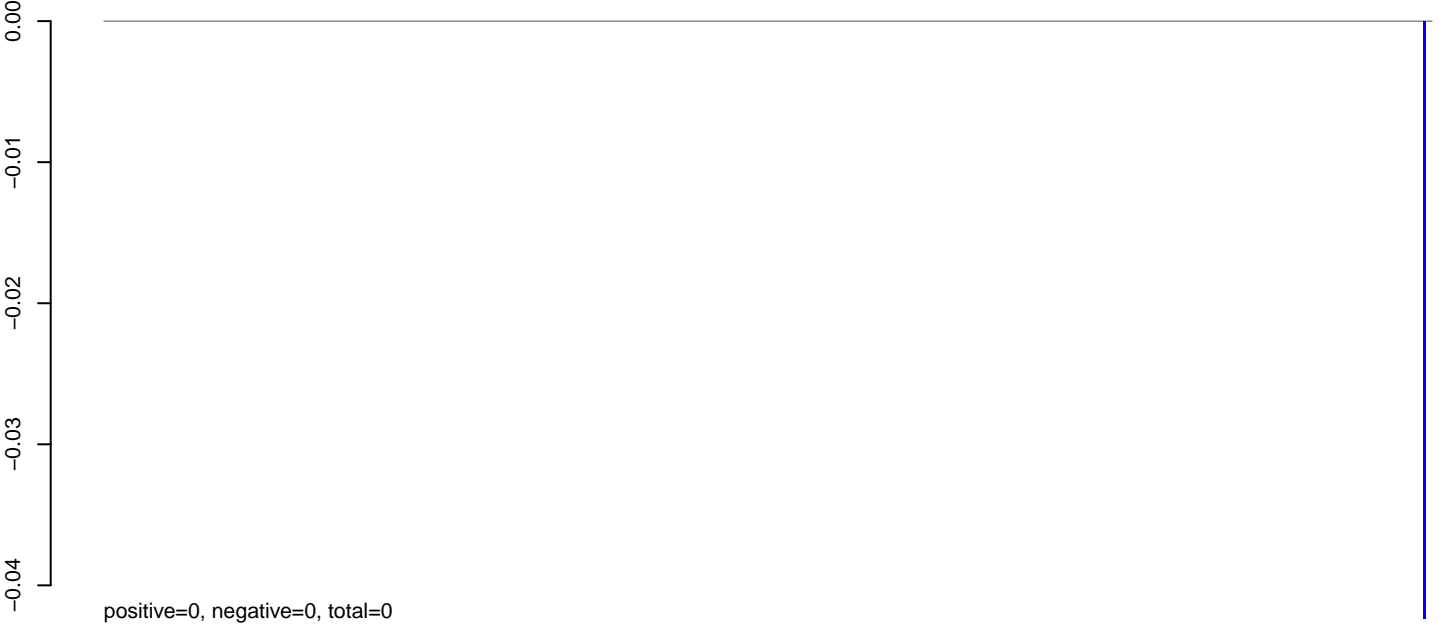
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



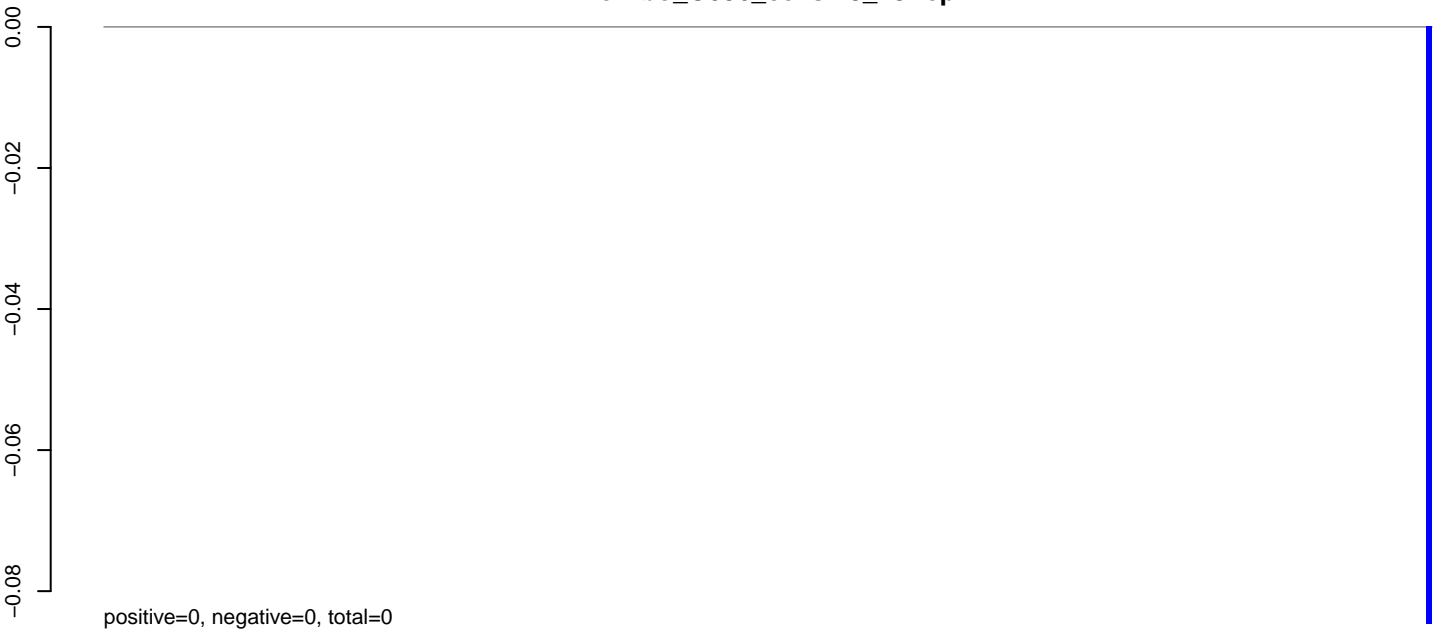
AeAlbo_C636_cells.24_35.rep



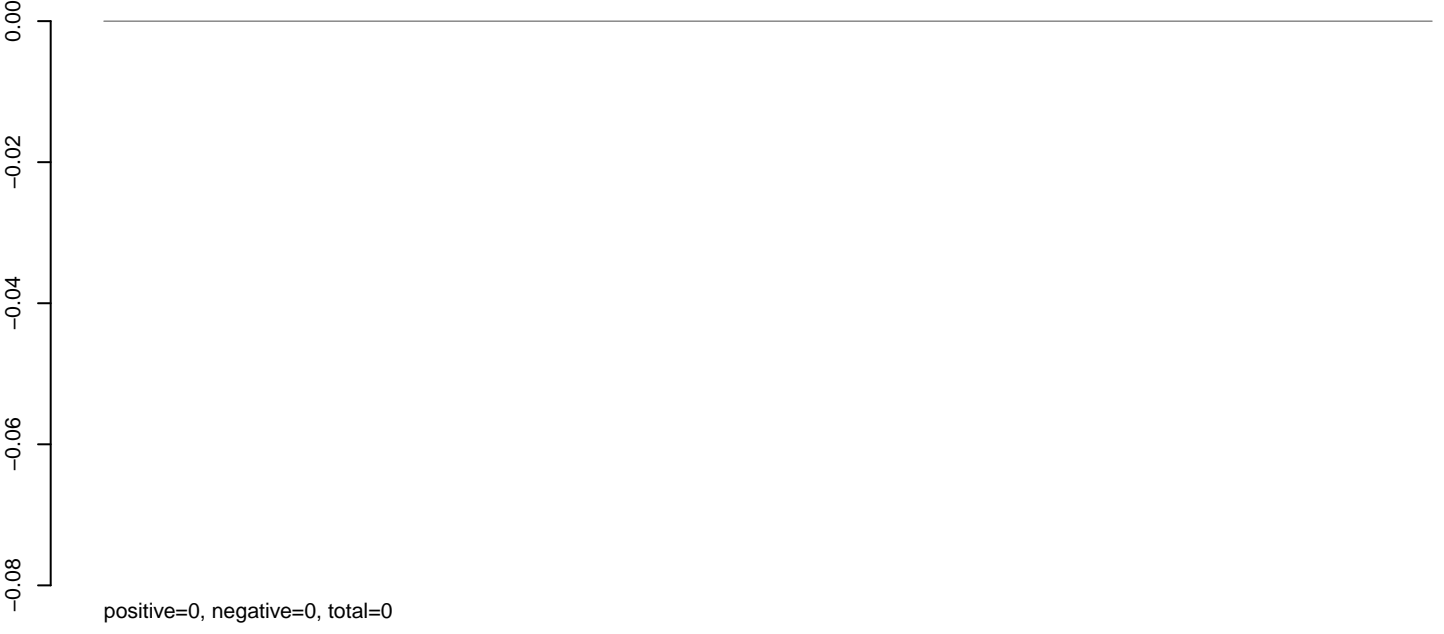
AeAlbo_C636_cells.rep



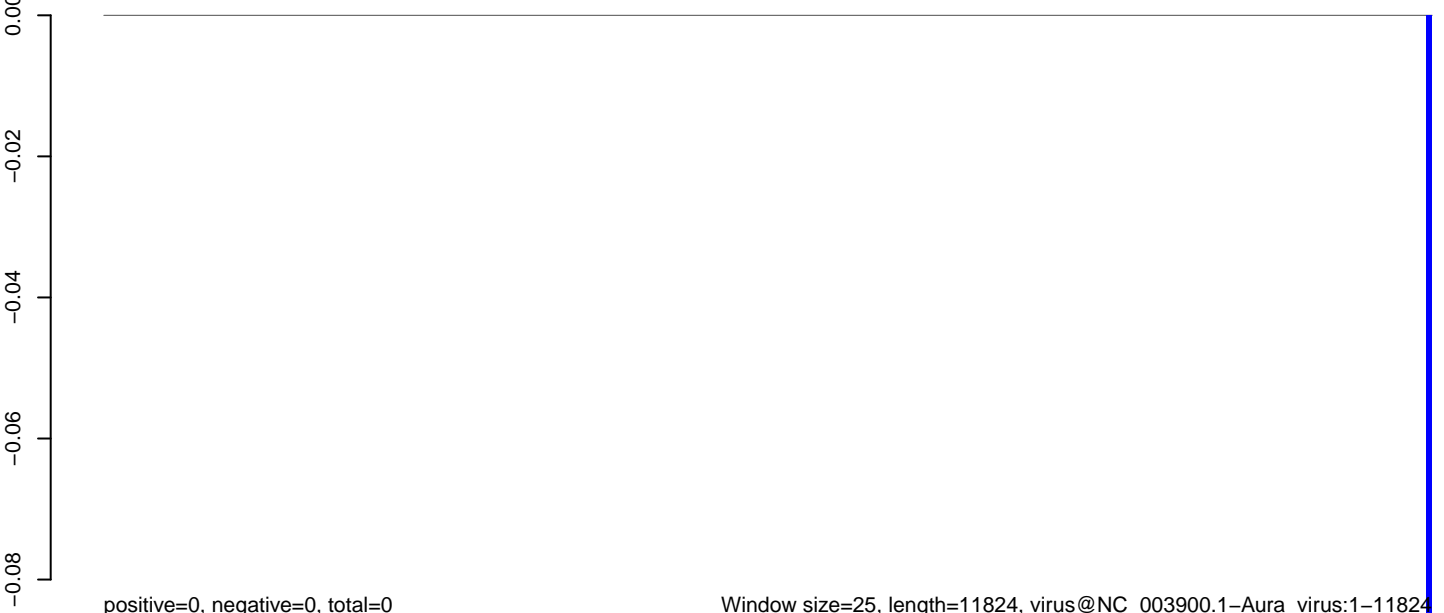
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



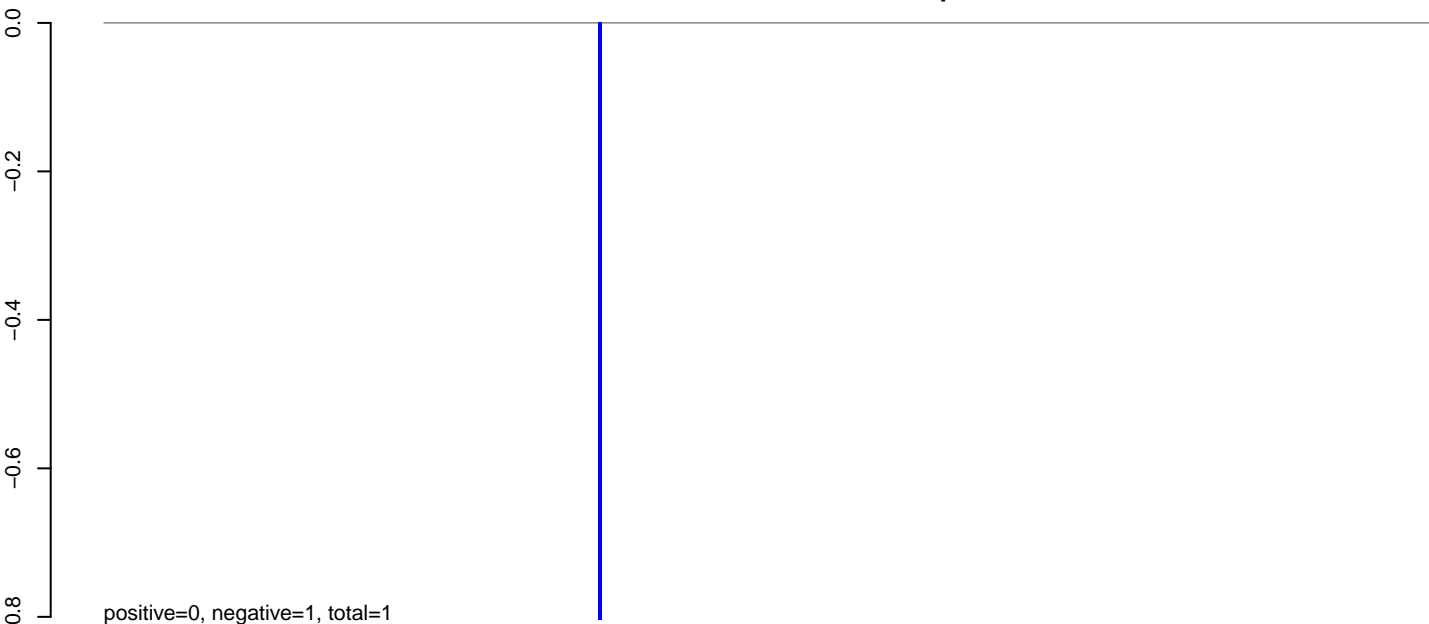
AeAlbo_C636_cells.rep



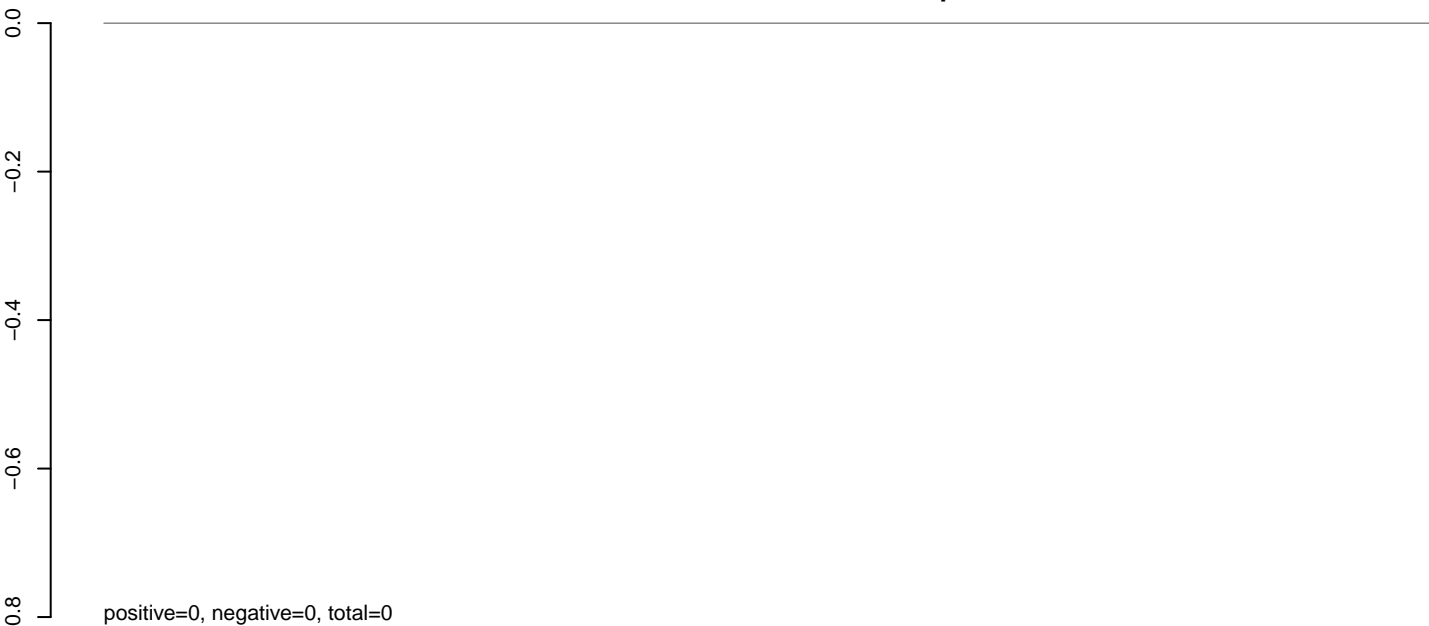
Window size=25, length=11824, virus@NC_003900.1-Aura_virus:1-11824

0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



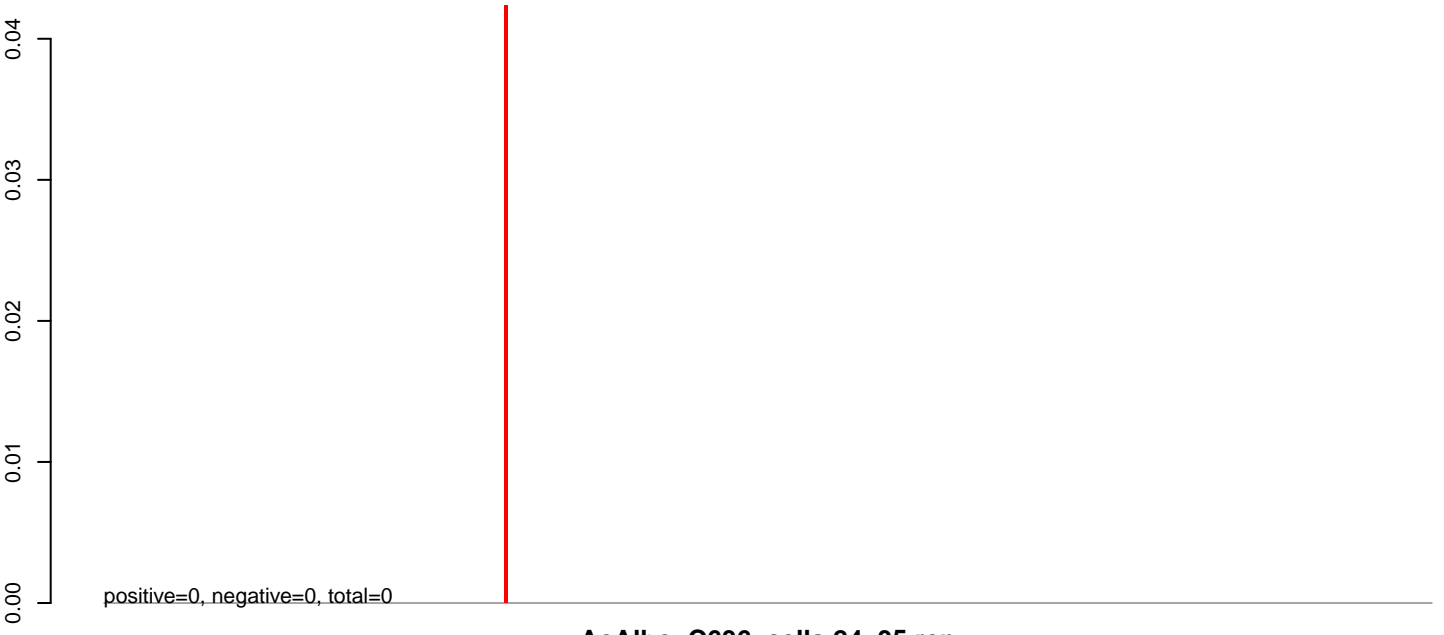
AeAlbo_C636_cells.rep



Window size=25, length=10943, virus@NC_003690.1-Langat_virus:1-10943

0 2000 4000 6000 8000 10000

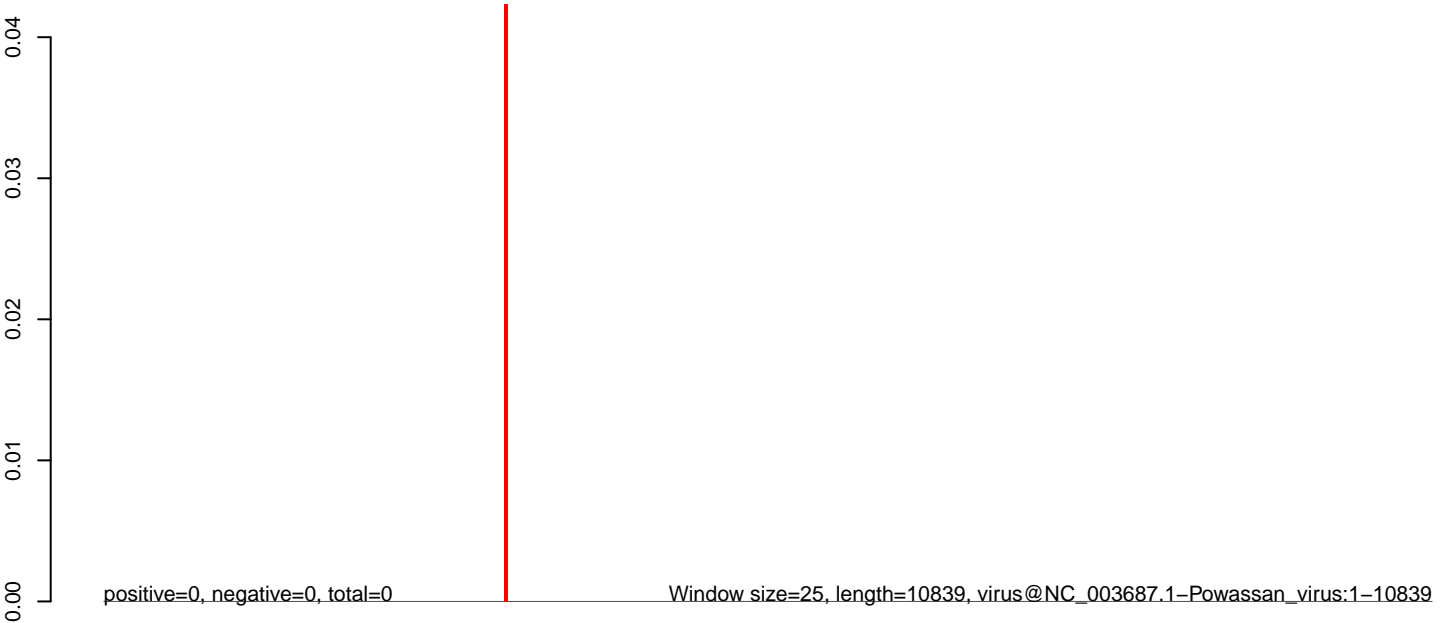
AeAlbo_C636_cells.18_23.rep



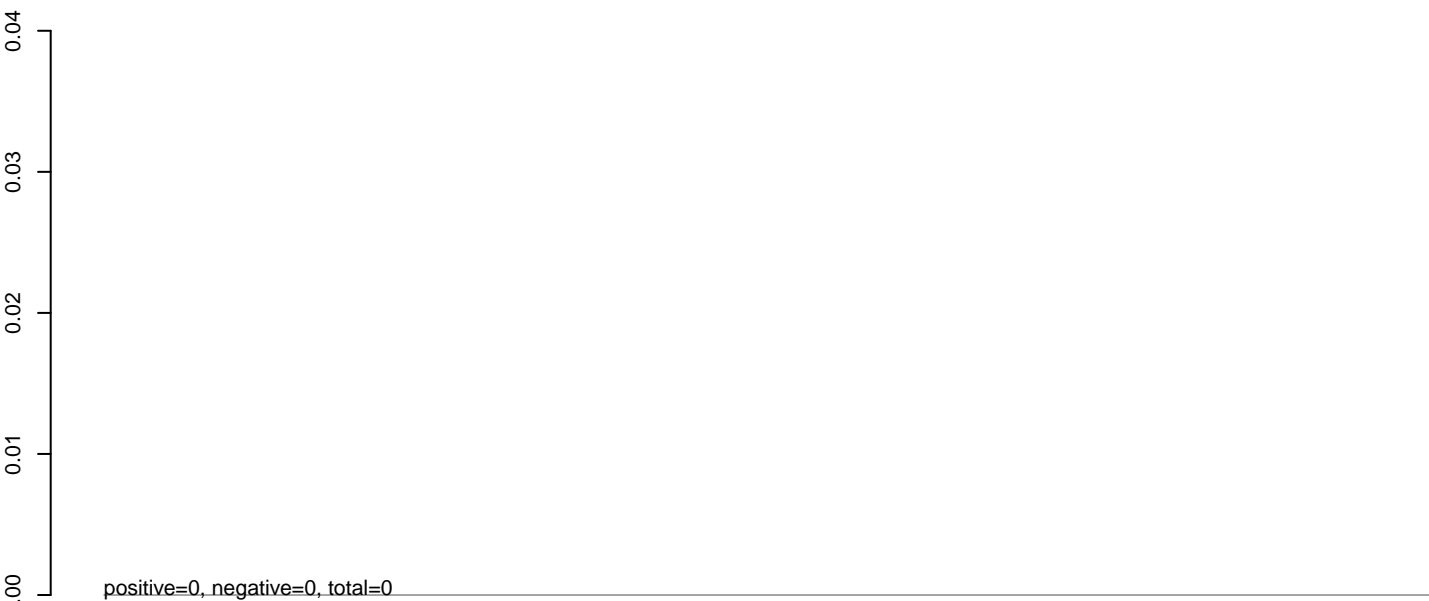
AeAlbo_C636_cells.24_35.rep



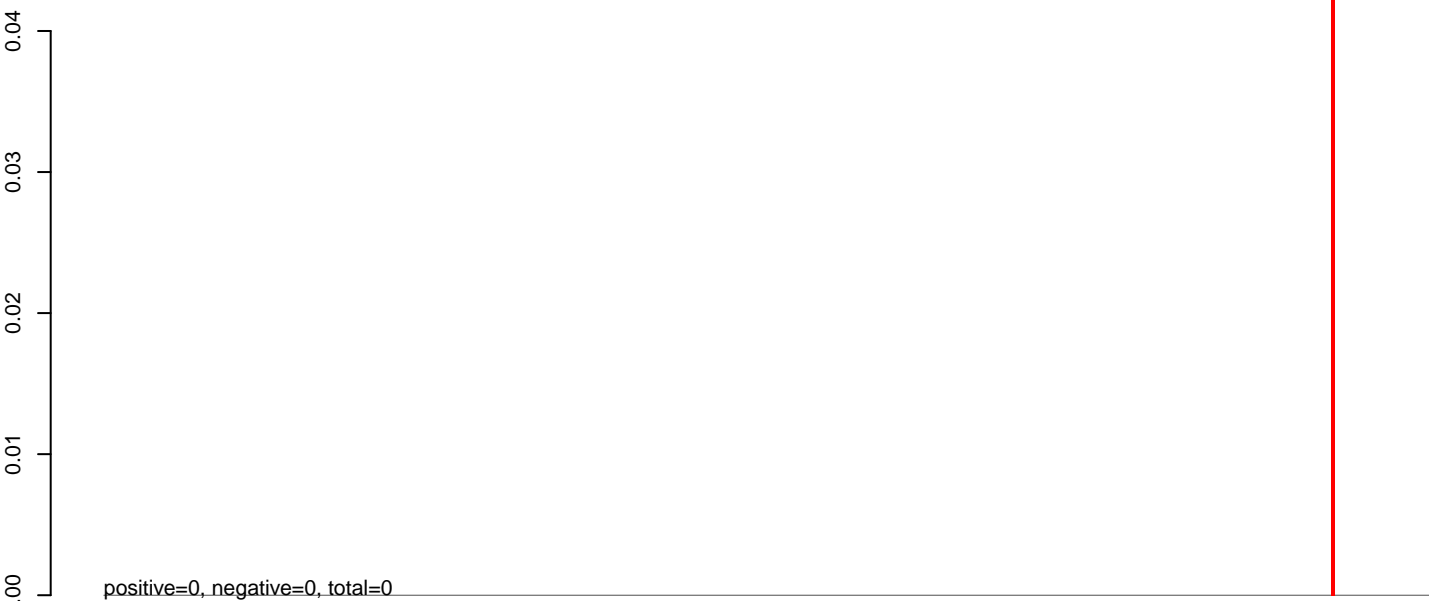
AeAlbo_C636_cells.rep



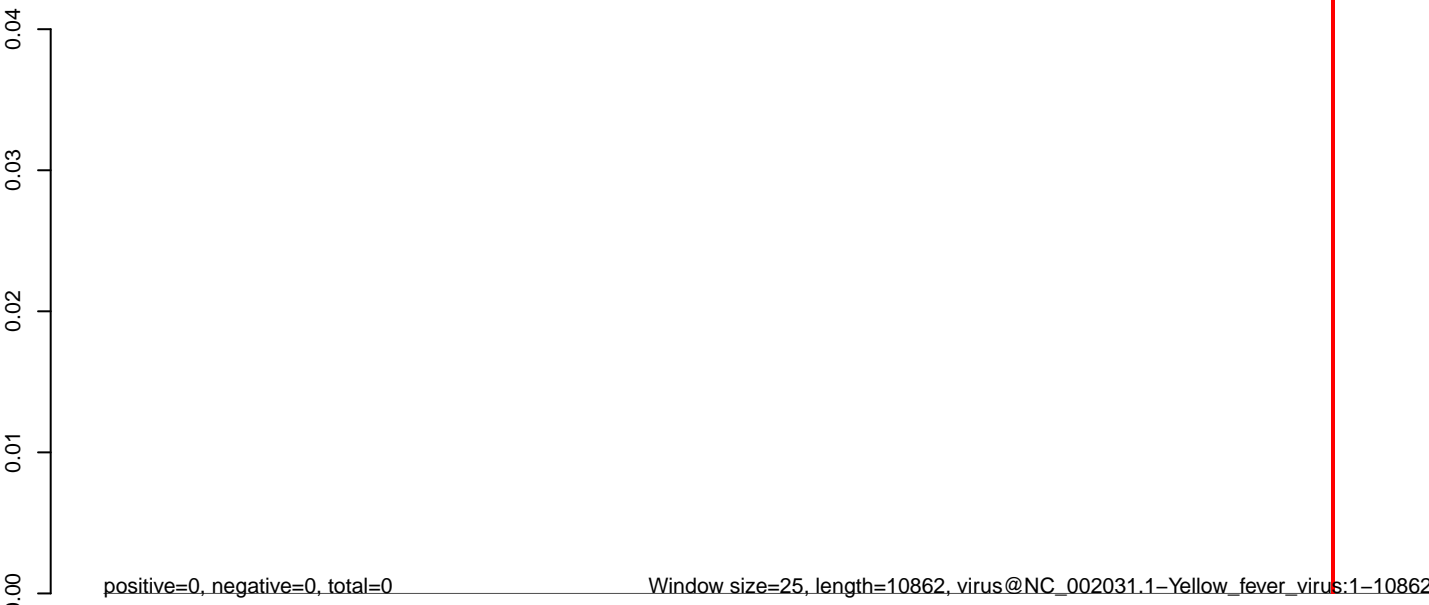
AeAlbo_C636_cells.18_23.rep



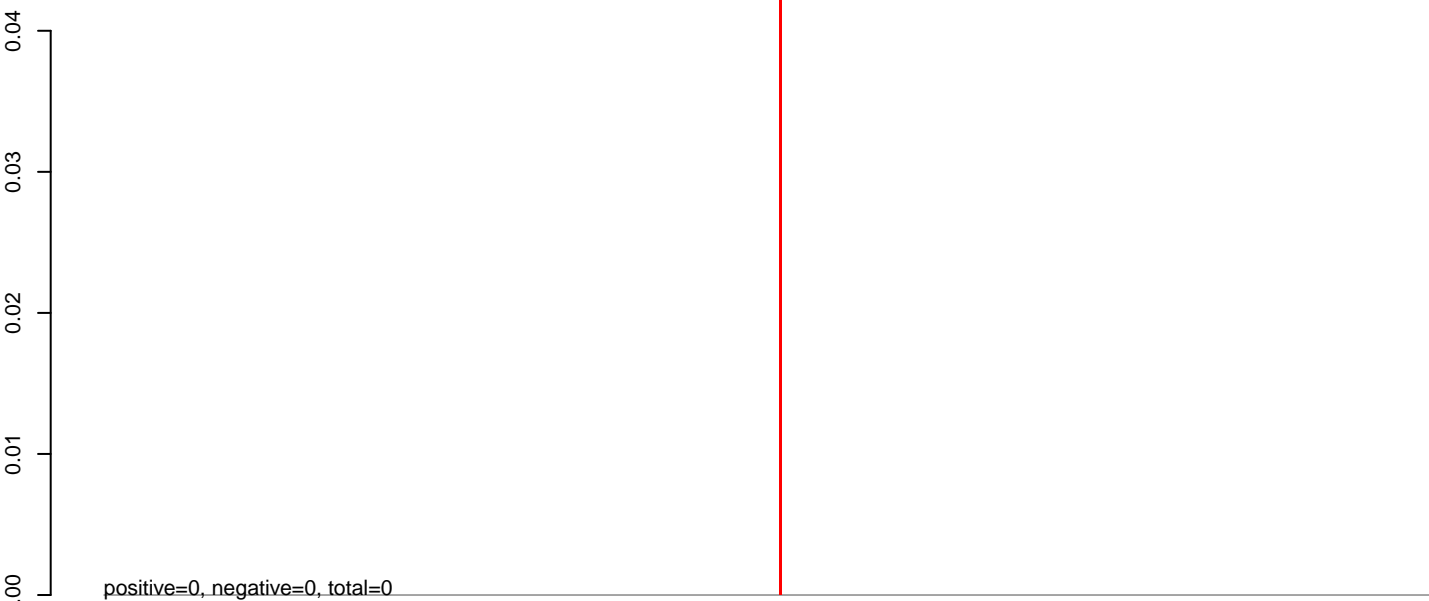
AeAlbo_C636_cells.24_35.rep



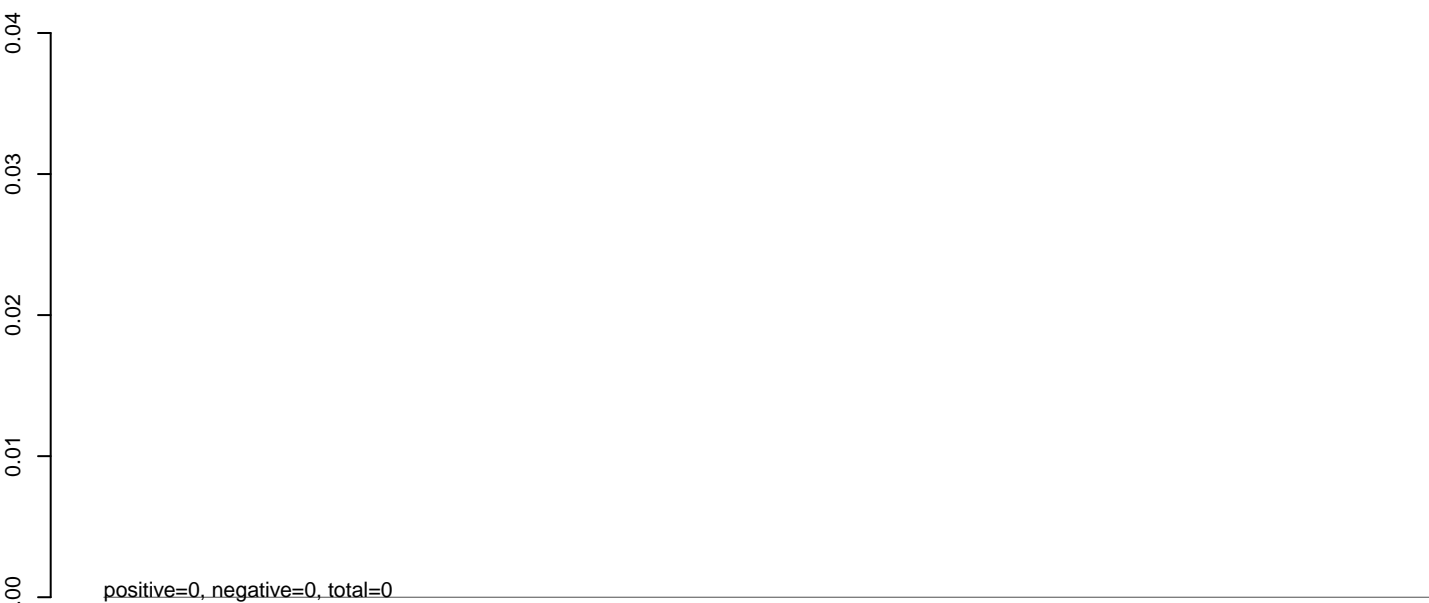
AeAlbo_C636_cells.rep



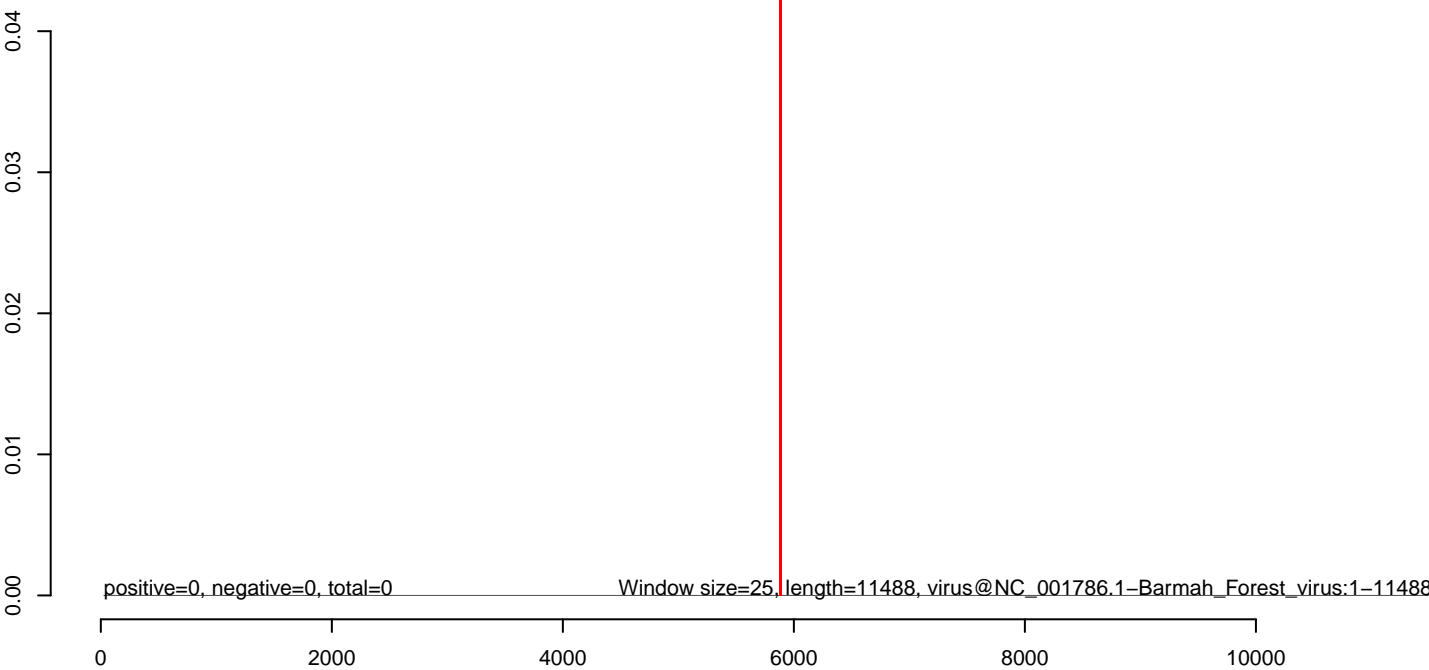
AeAlbo_C636_cells.18_23.rep



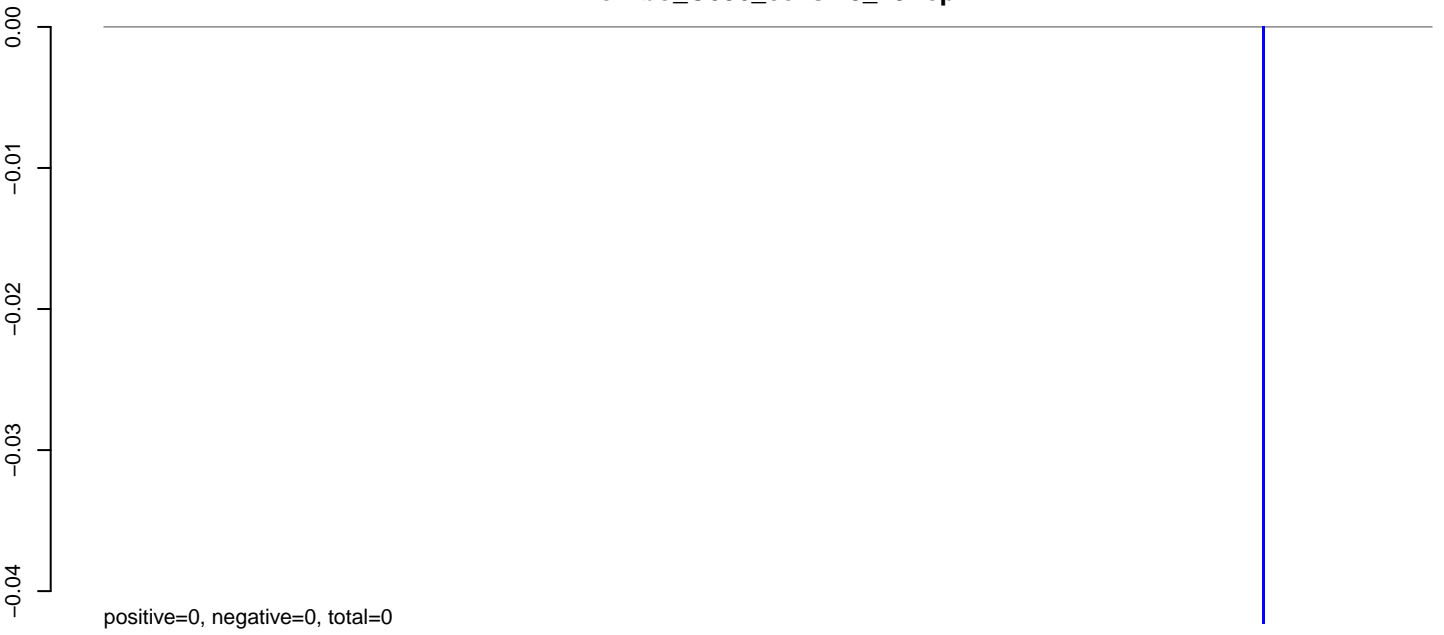
AeAlbo_C636_cells.24_35.rep



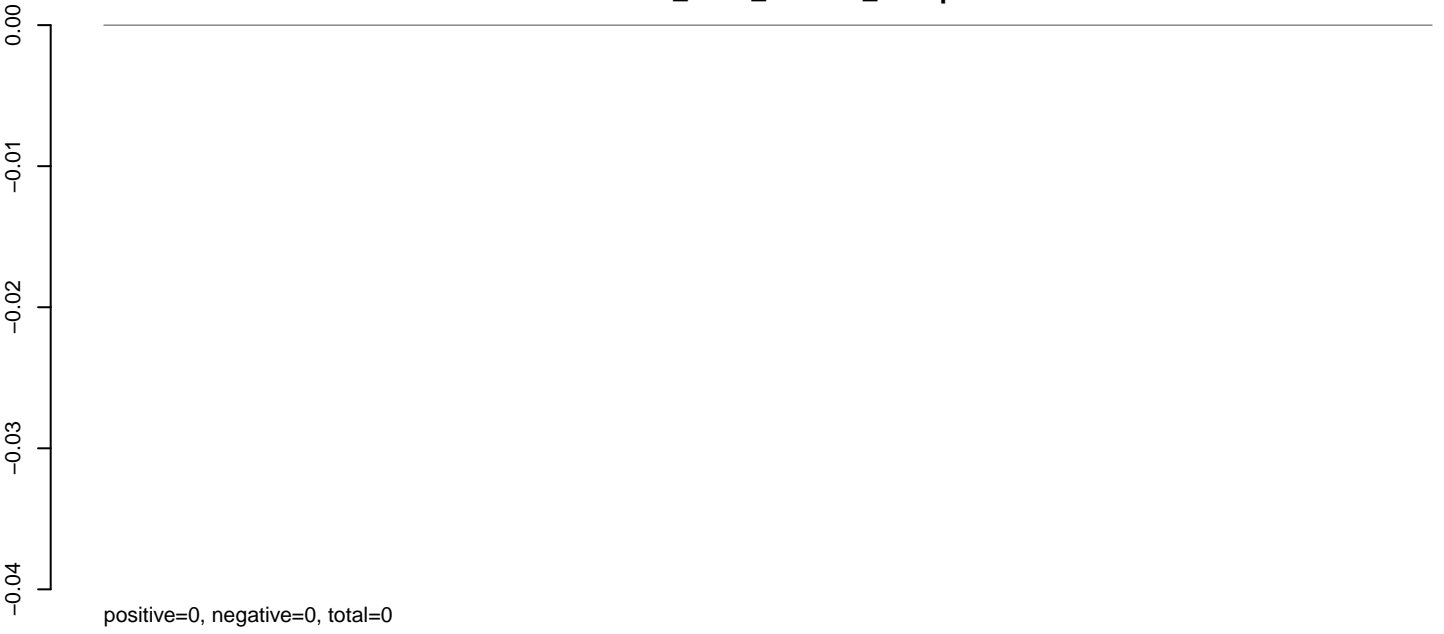
AeAlbo_C636_cells.rep



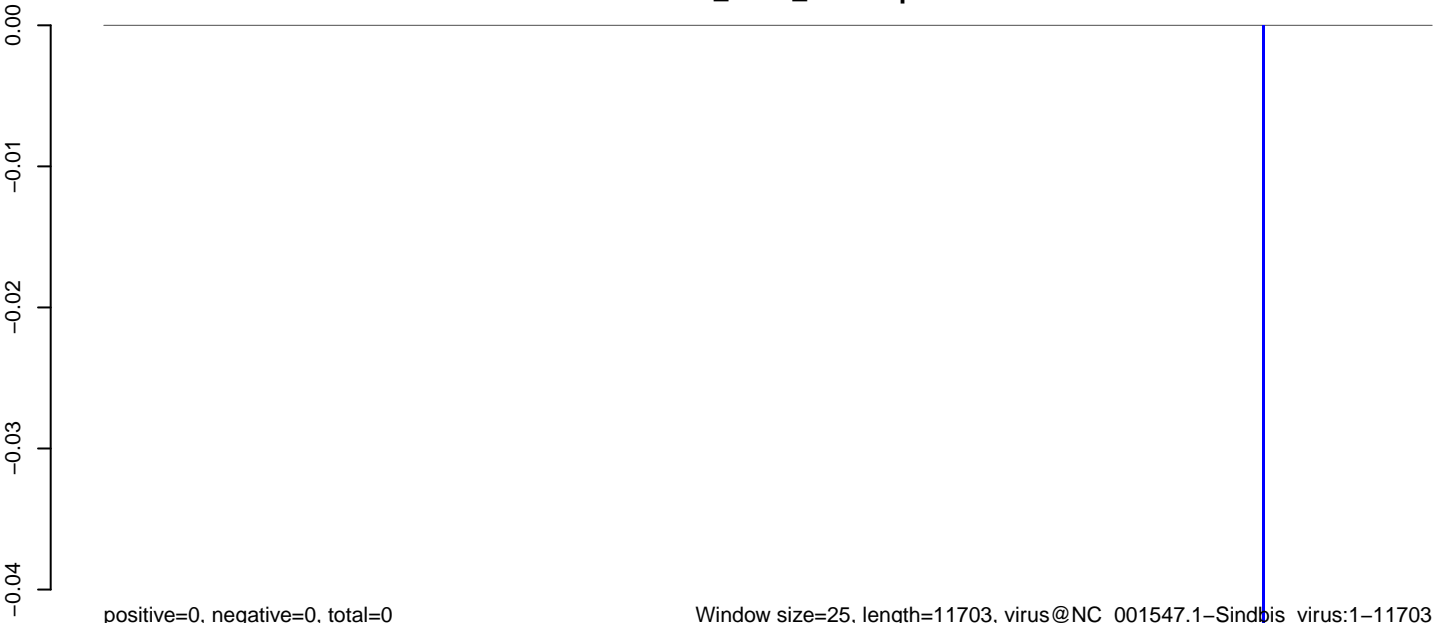
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



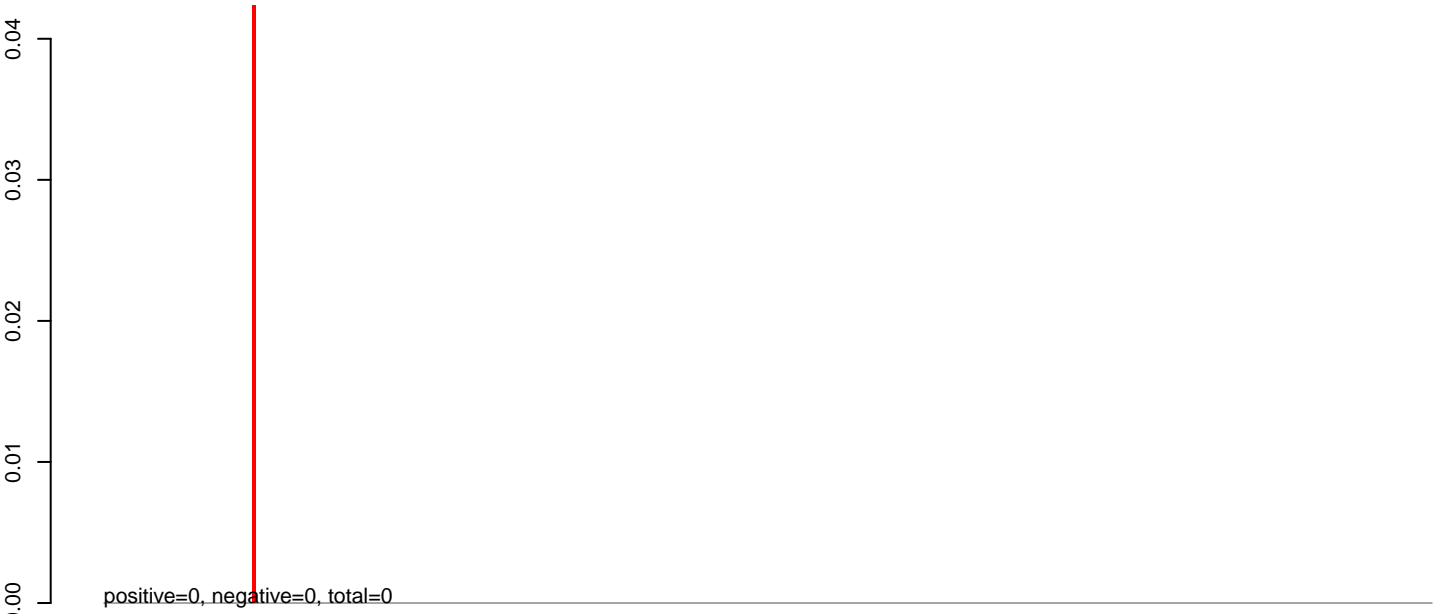
AeAlbo_C636_cells.rep



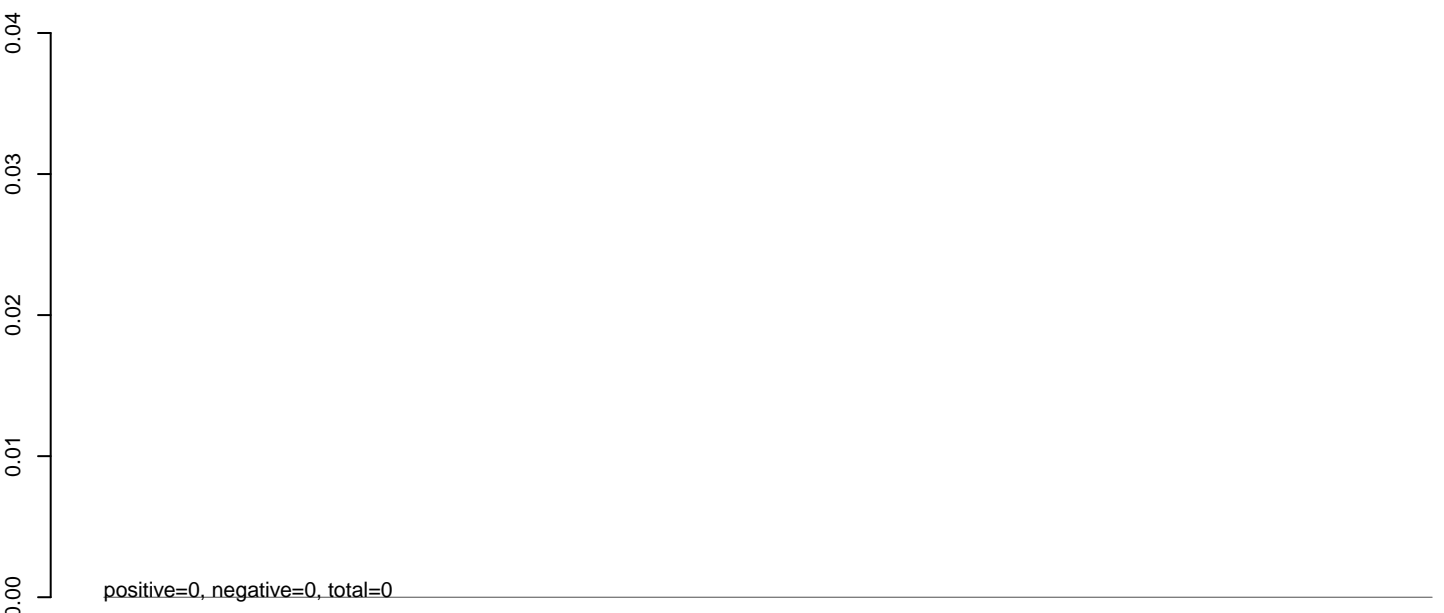
Window size=25, length=11703, virus@NC_001547.1-Sindbis_virus:1-11703

0 2000 4000 6000 8000 10000 12000

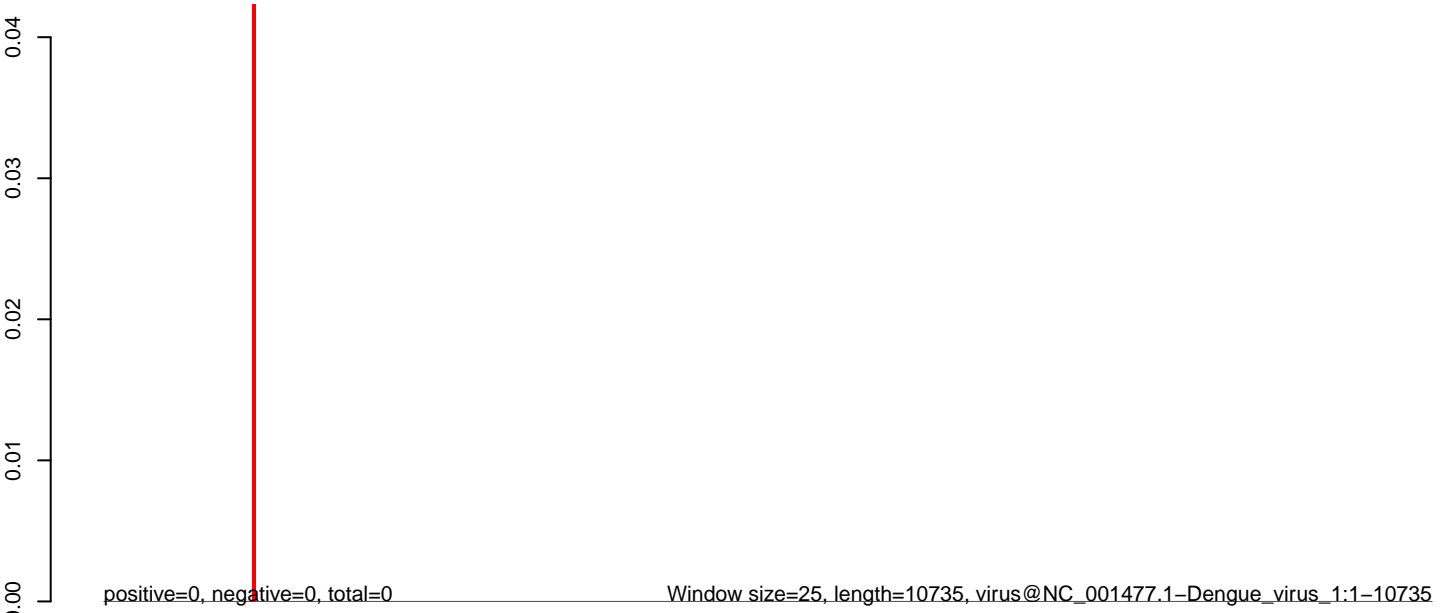
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

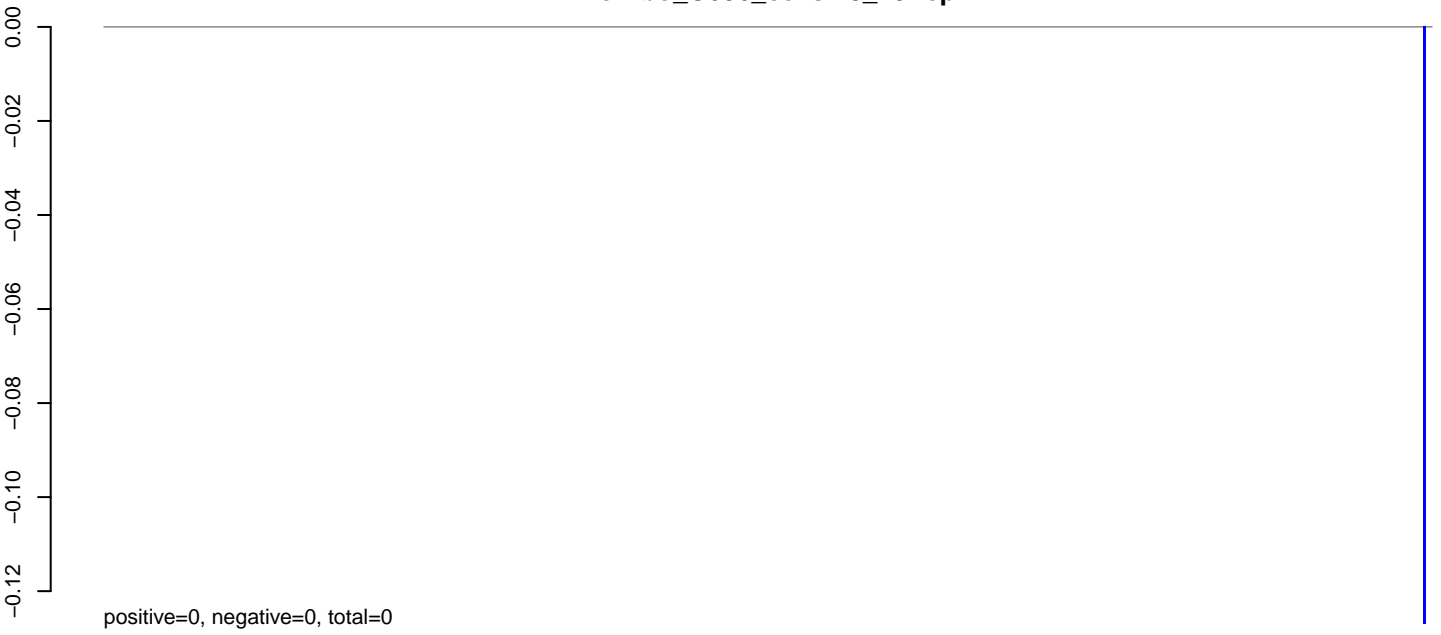


AeAlbo_C636_cells.rep

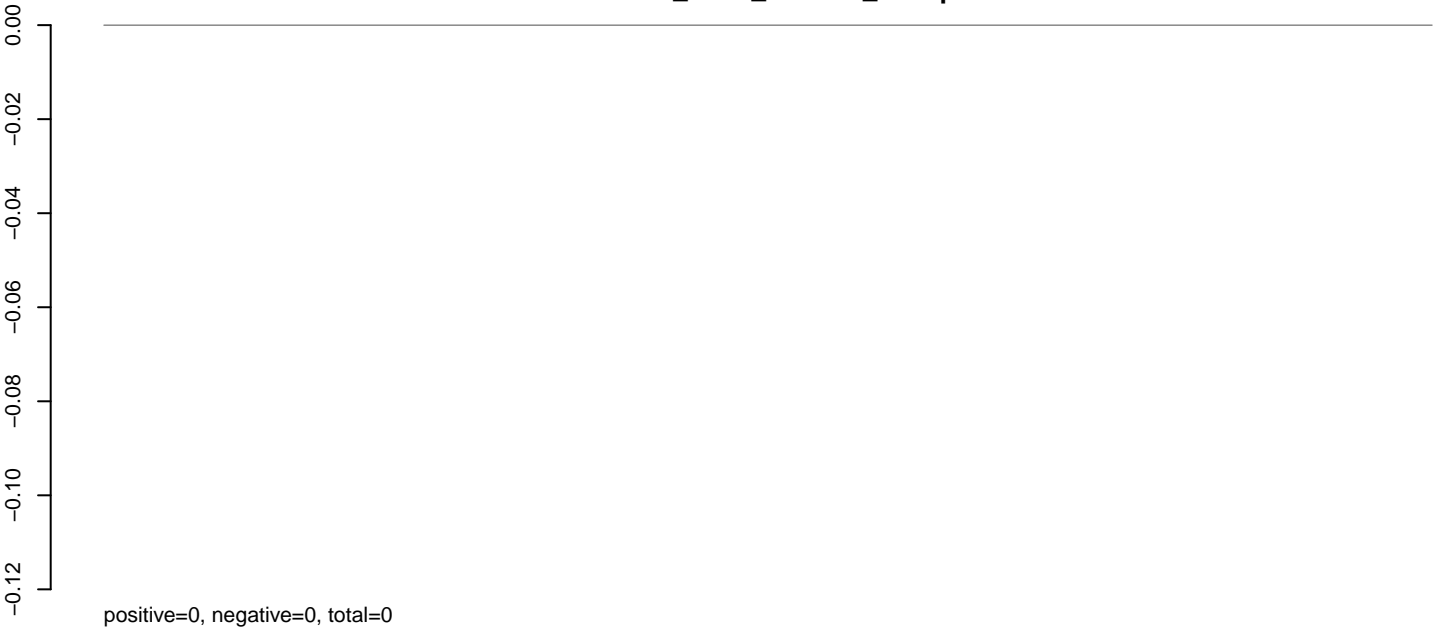


Window size=25, length=10735, virus@NC_001477.1-Dengue_virus_1:1-10735

AeAlbo_C636_cells.18_23.rep



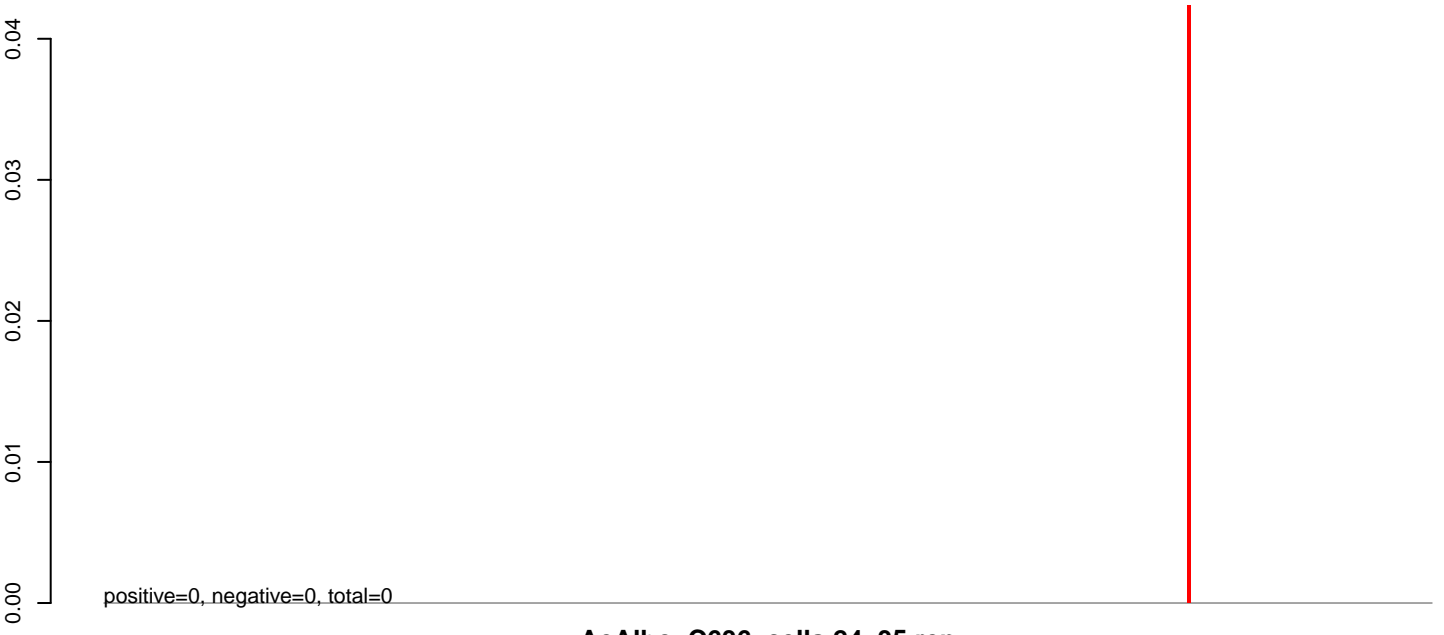
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



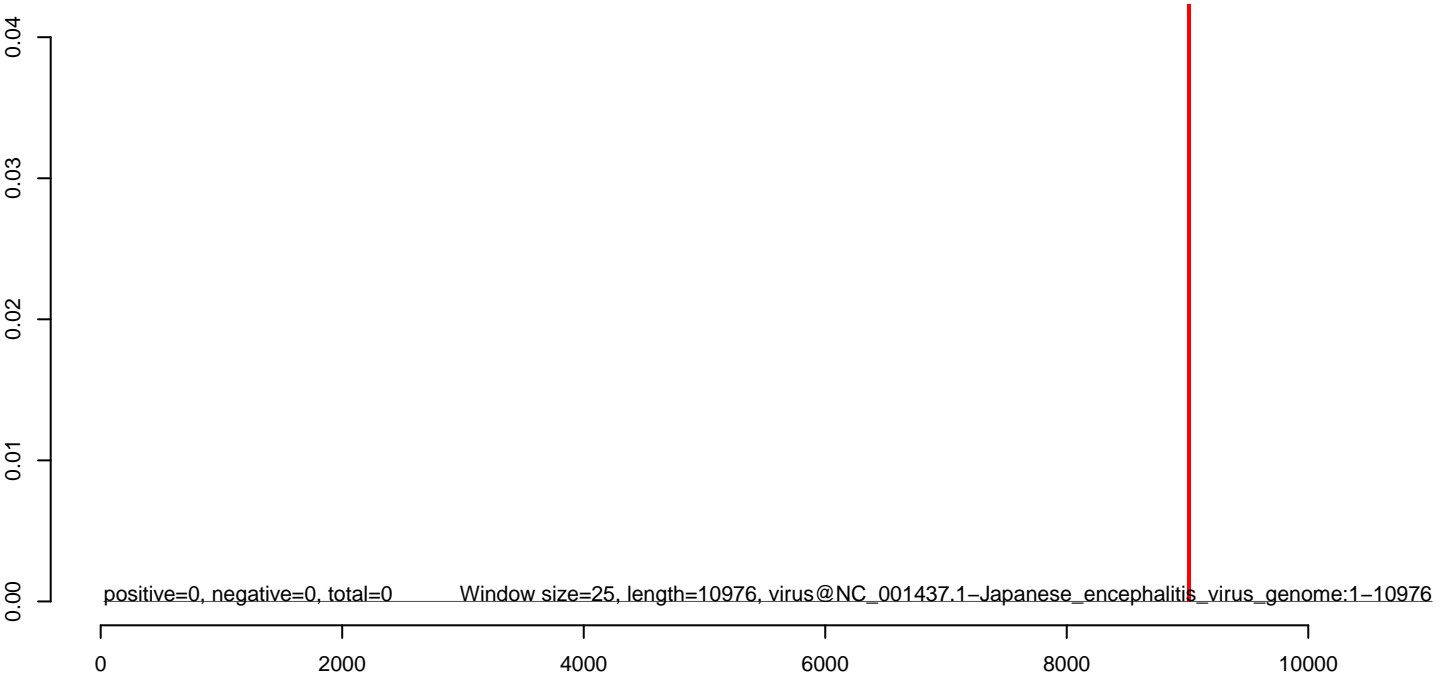
AeAlbo_C636_cells.18_23.rep



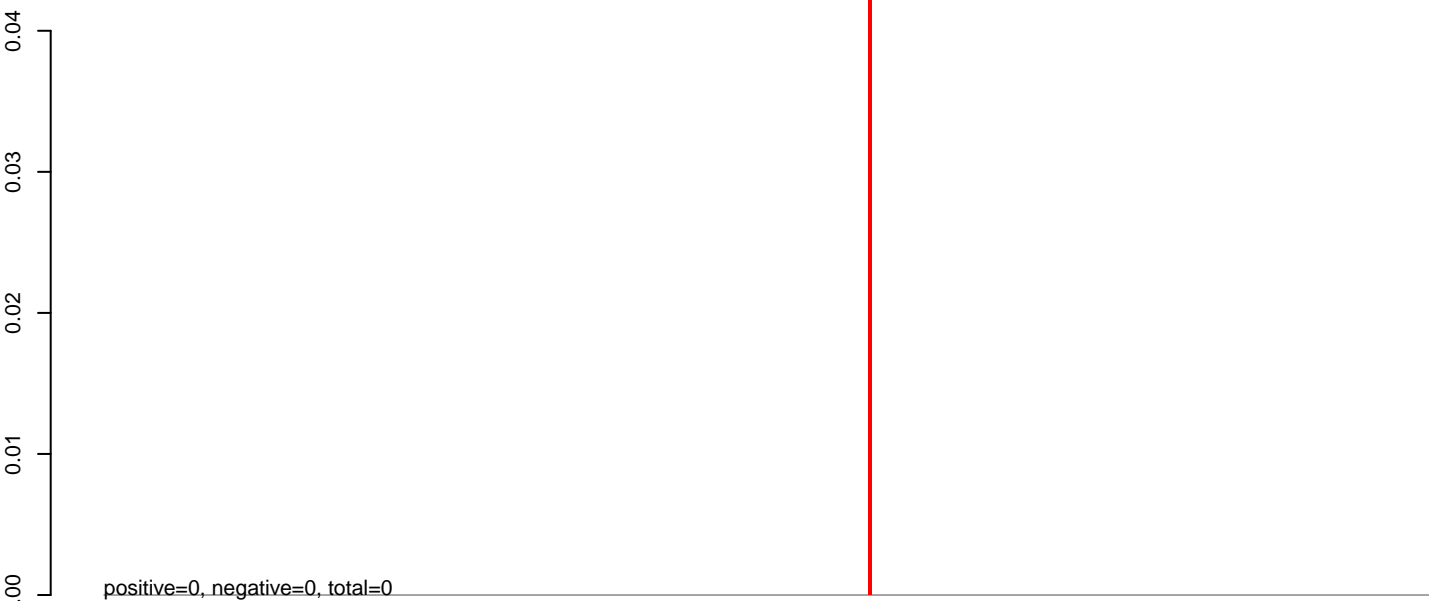
AeAlbo_C636_cells.24_35.rep



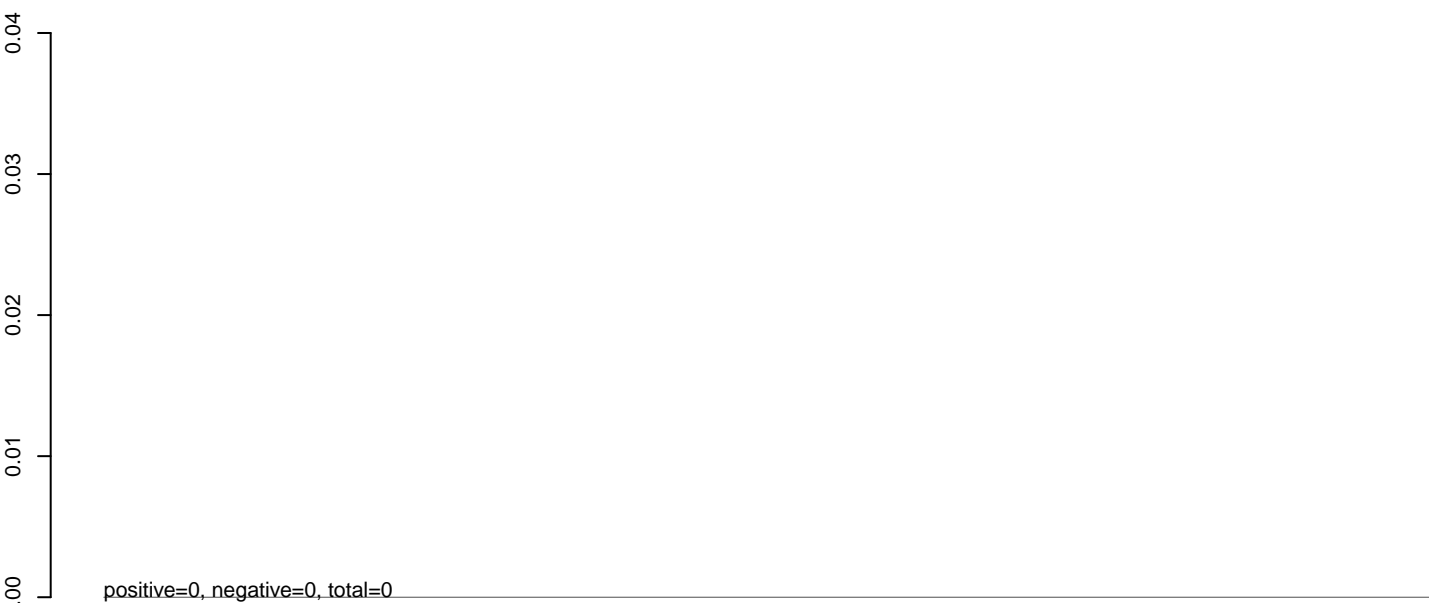
AeAlbo_C636_cells.rep



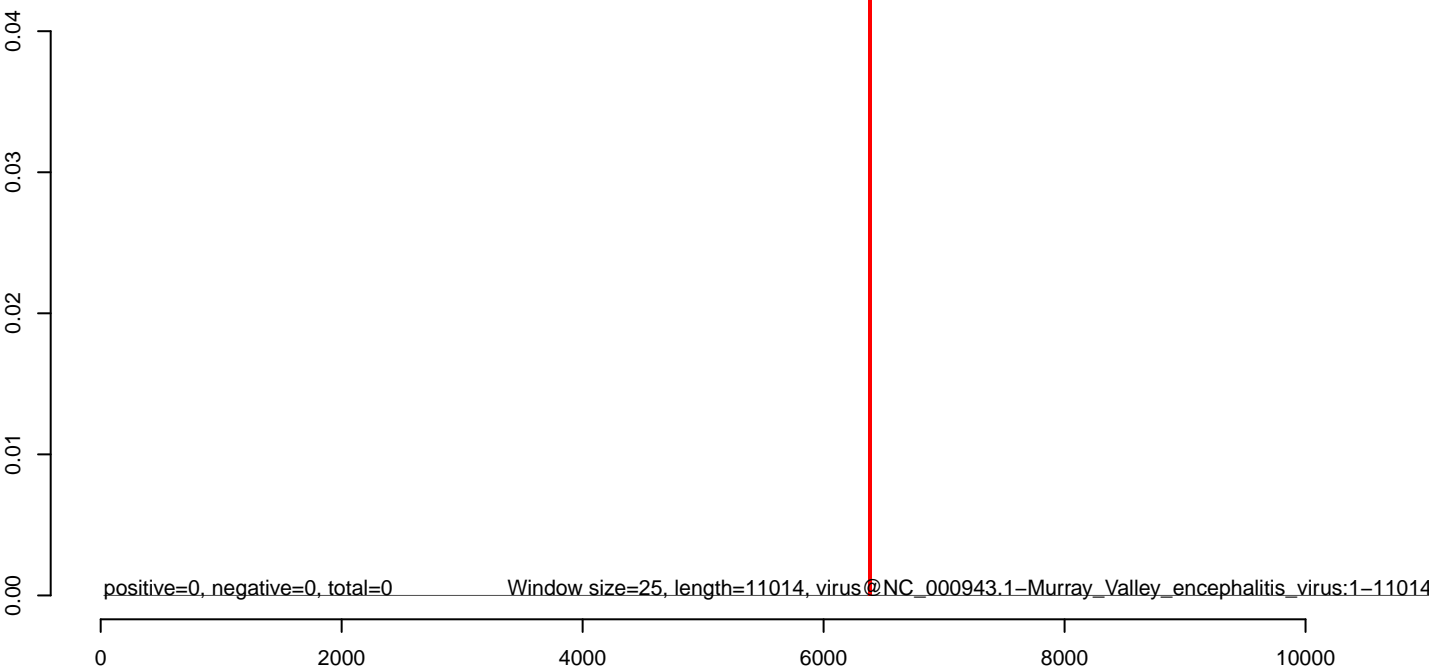
AeAlbo_C636_cells.18_23.rep



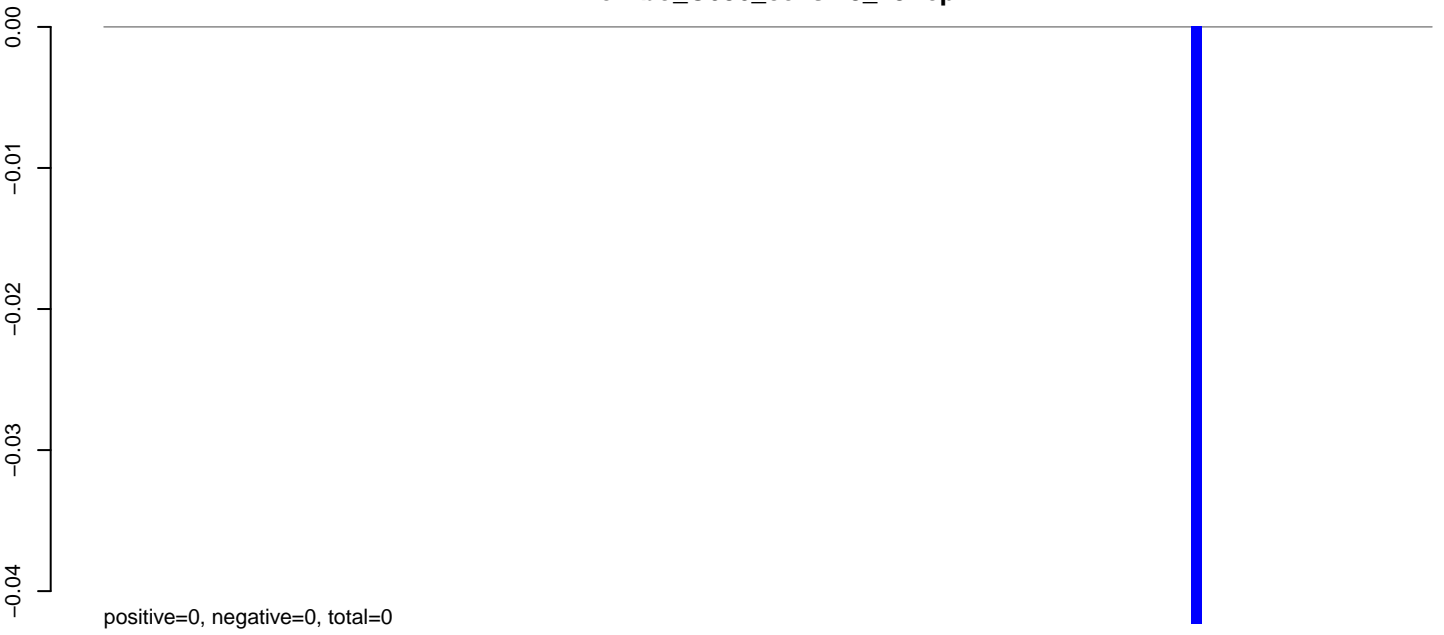
AeAlbo_C636_cells.24_35.rep



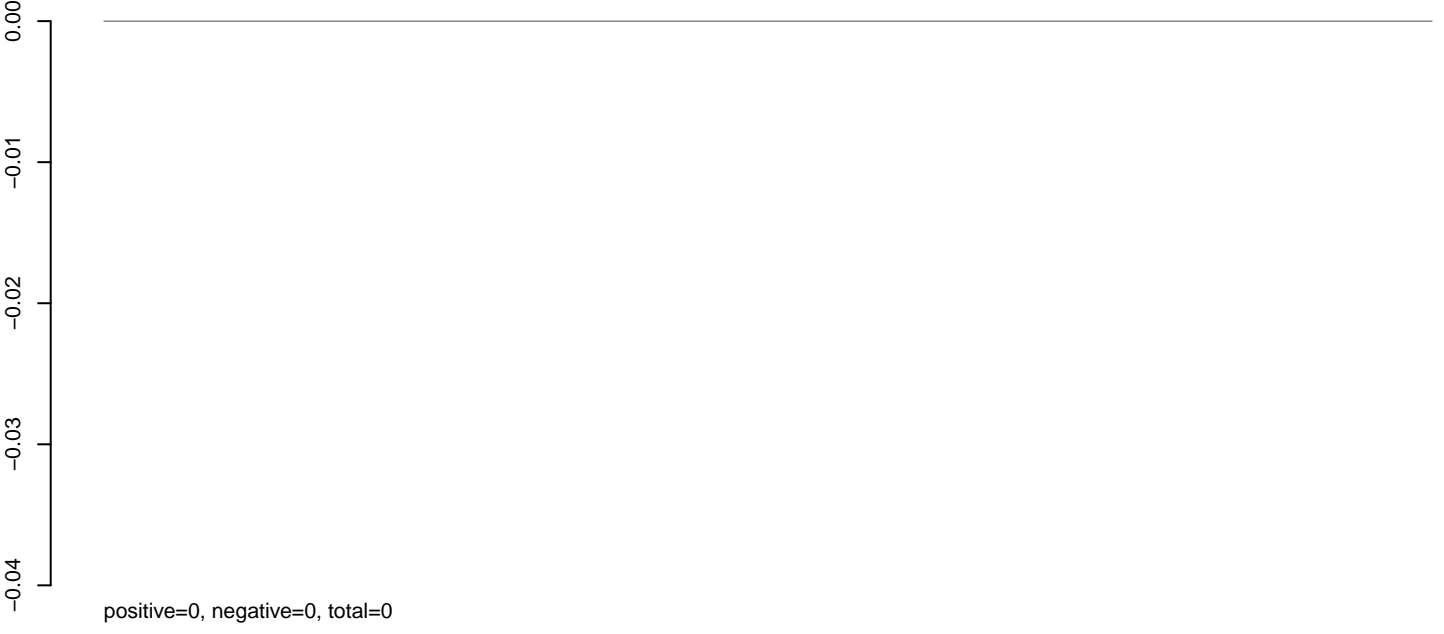
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



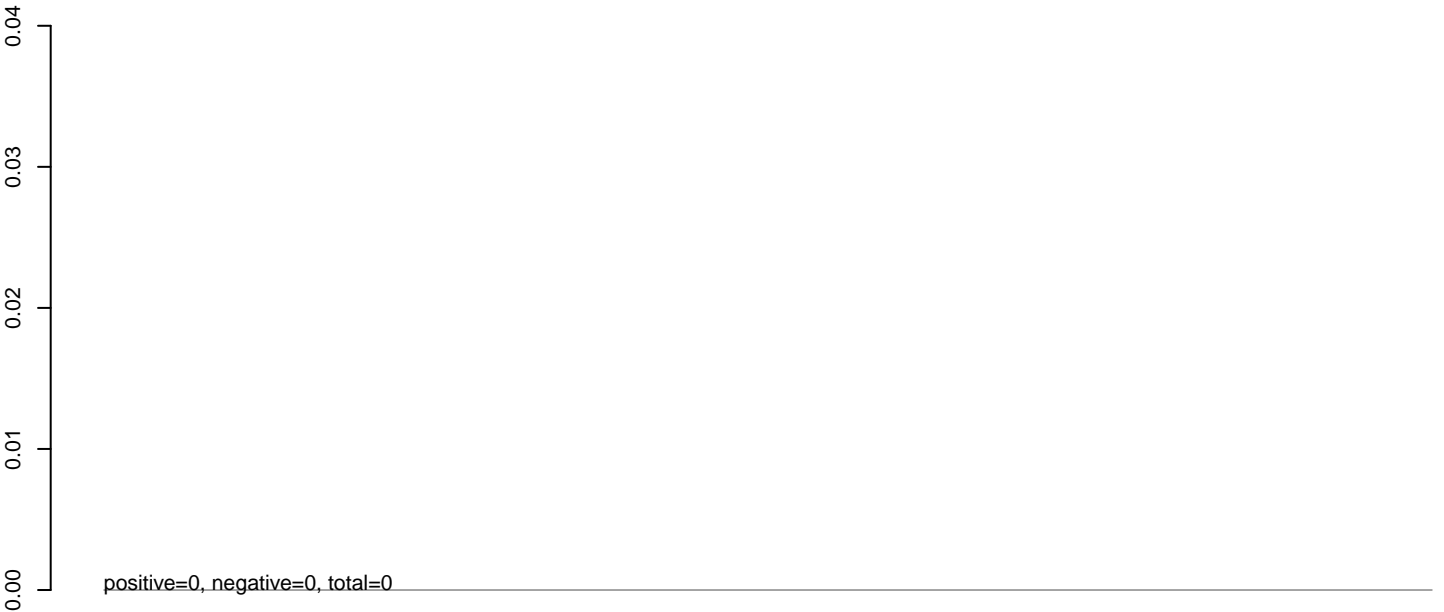
AeAlbo_C636_cells.rep



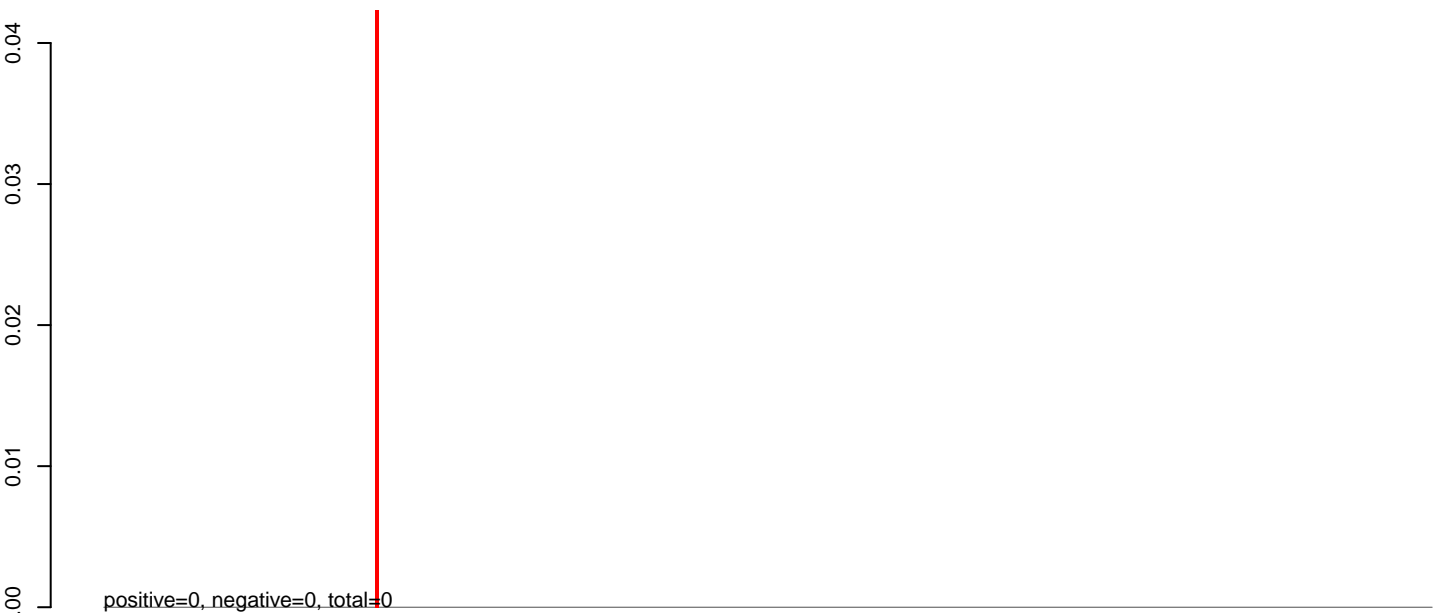
Window size=25, length=3157, virus@MH188043.1-Culex_densovirus_strain_CDENV_Madera-2:1-3157

0 500 1000 1500 2000 2500 3000

AeAlbo_C636_cells.18_23.rep



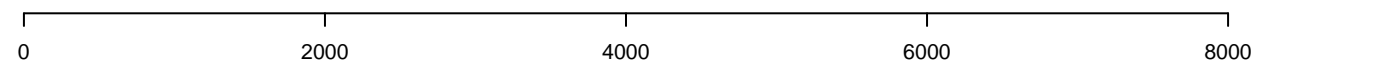
AeAlbo_C636_cells.24_35.rep



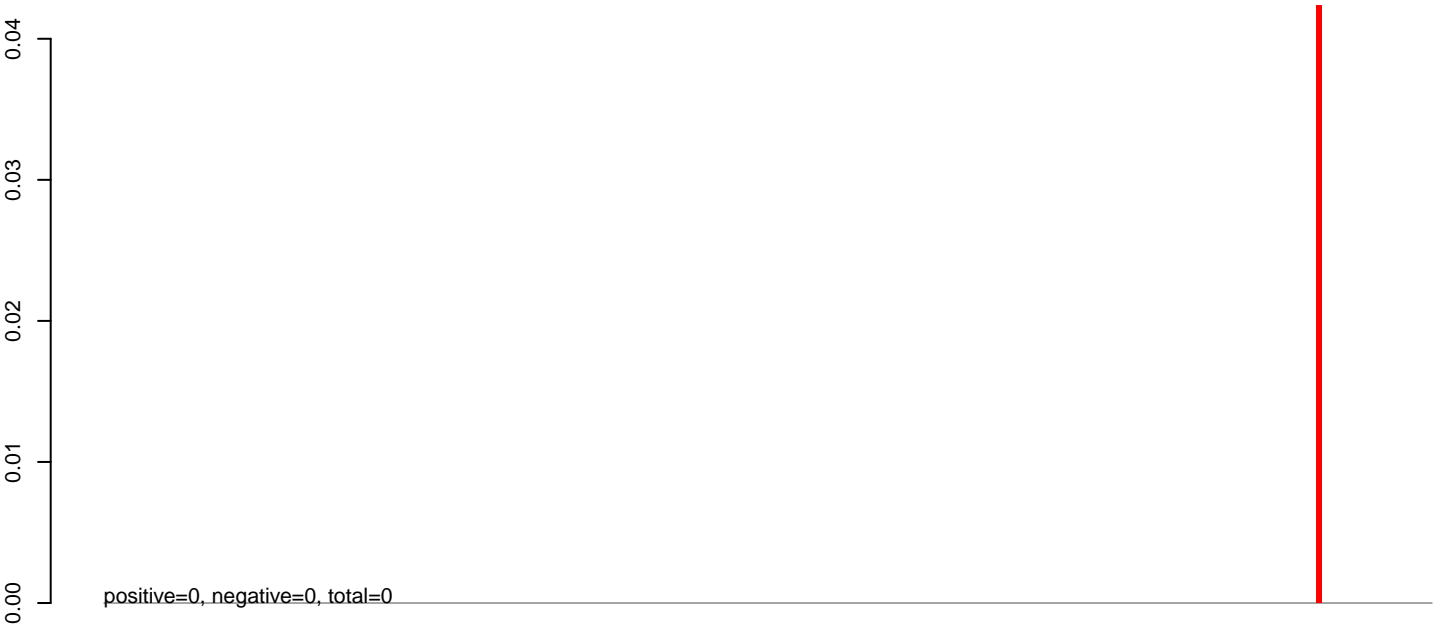
AeAlbo_C636_cells.rep



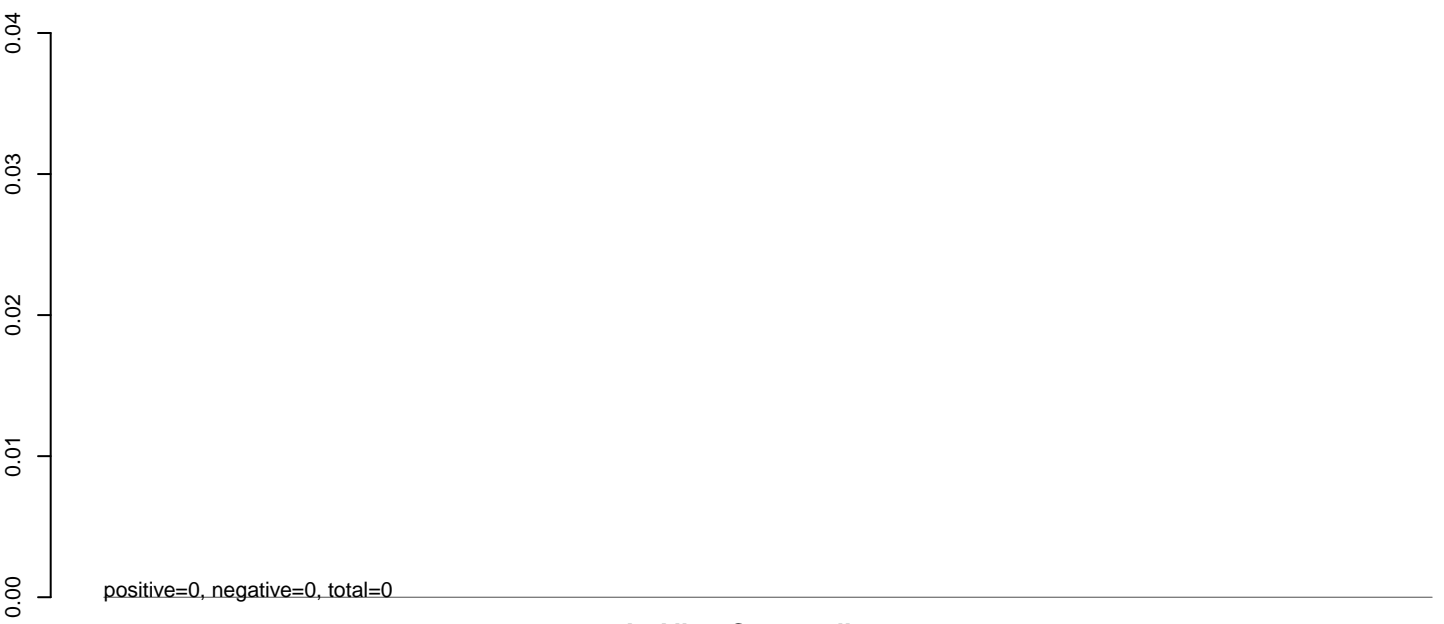
downloaded from SRA on 2025-10-14 10:19:03 AM. File: 176325.1_Aedes_alboannulatus_orthomyxo-like_virus_strain_mosWSAb70818_segment_1_2_3_4:1-8819



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

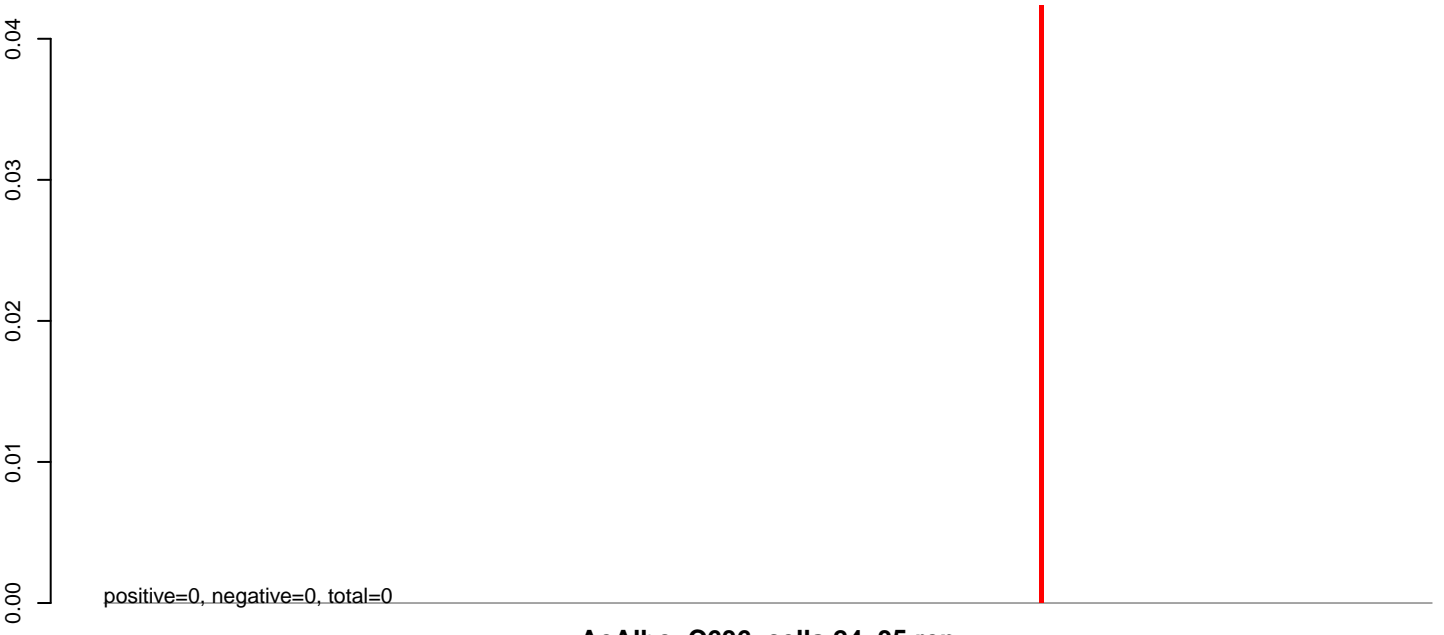


AeAlbo_C636_cells.rep



negative-size=25, length=5982, virus@MF176317.1-Culex_mononega-like_virus_1_strain_mos191X45127_segment_2:1-5982

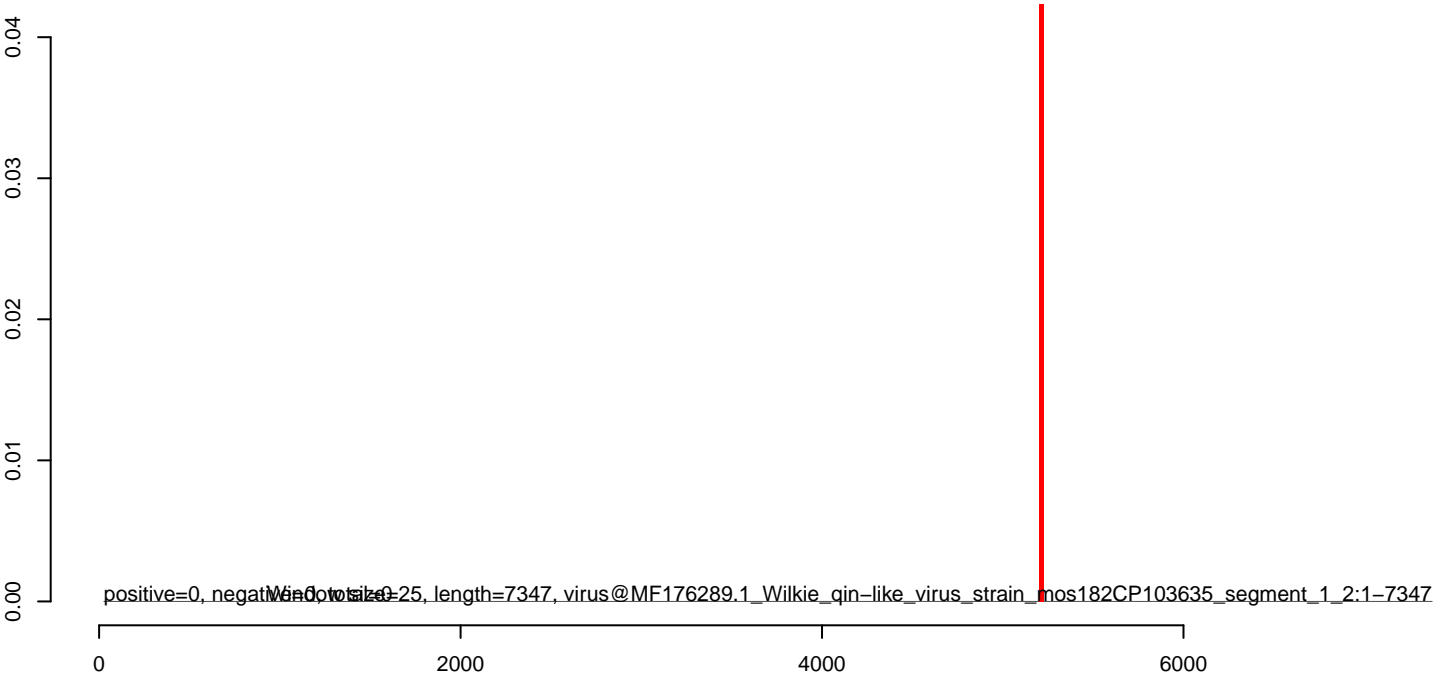
AeAlbo_C636_cells.18_23.rep



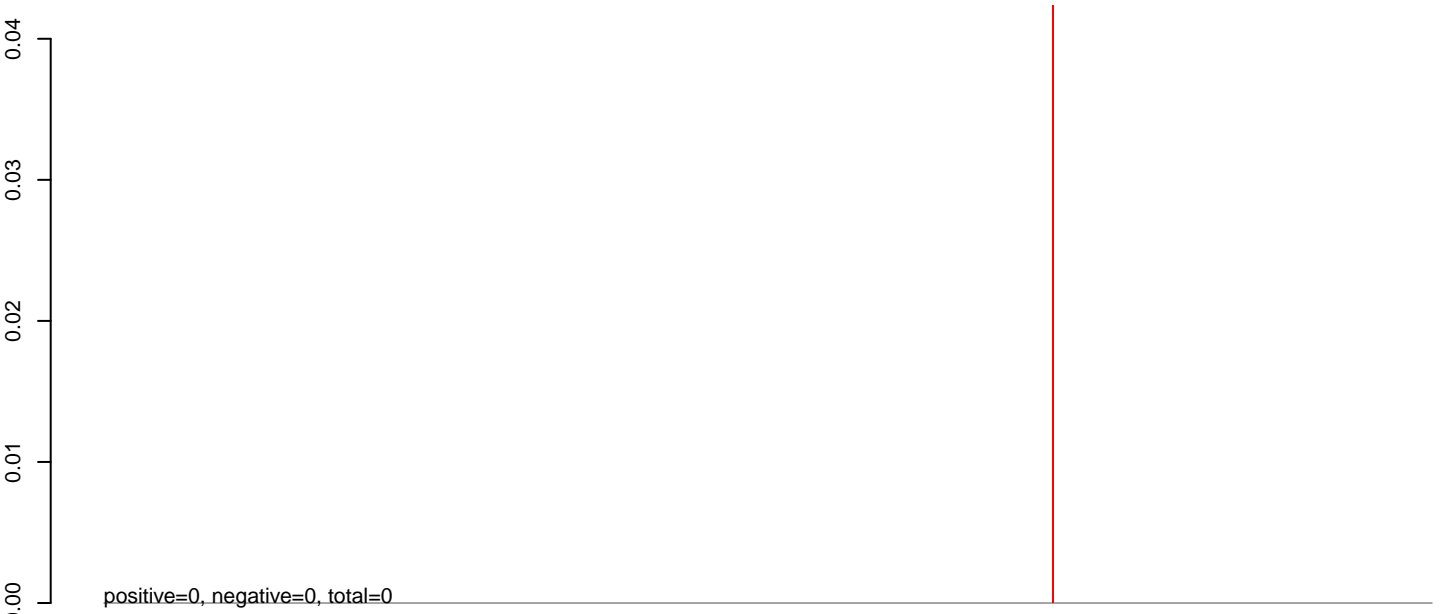
AeAlbo_C636_cells.24_35.rep



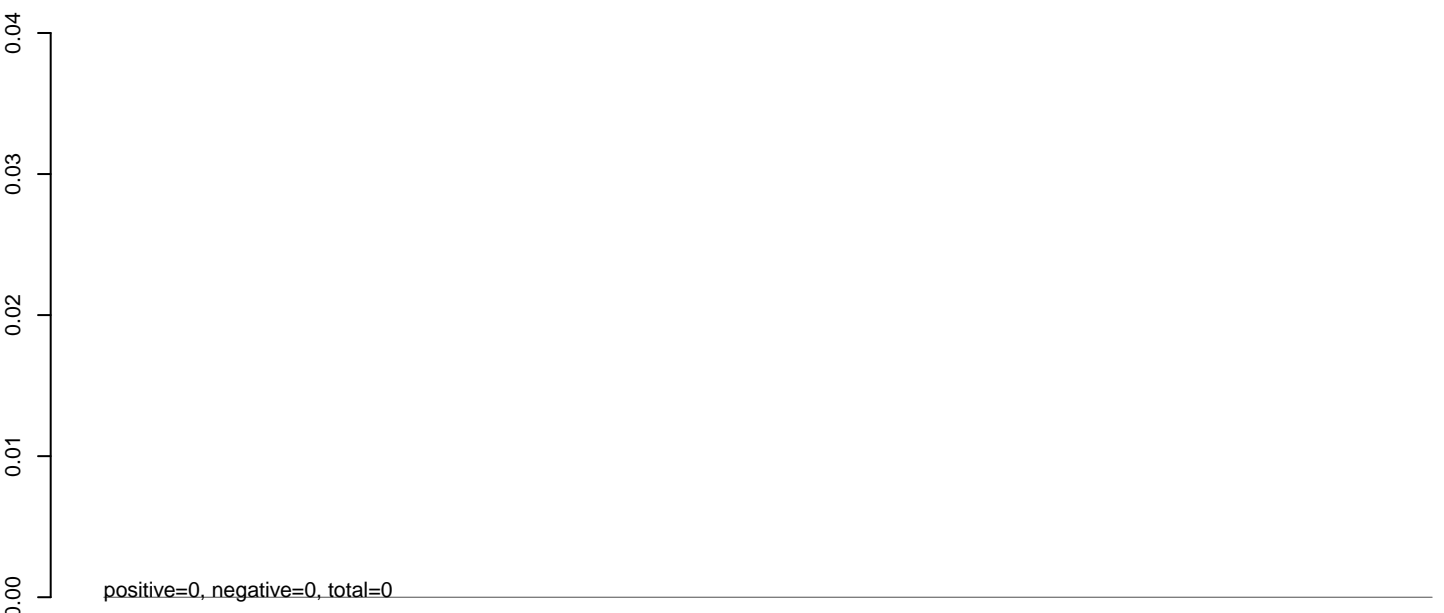
AeAlbo_C636_cells.rep



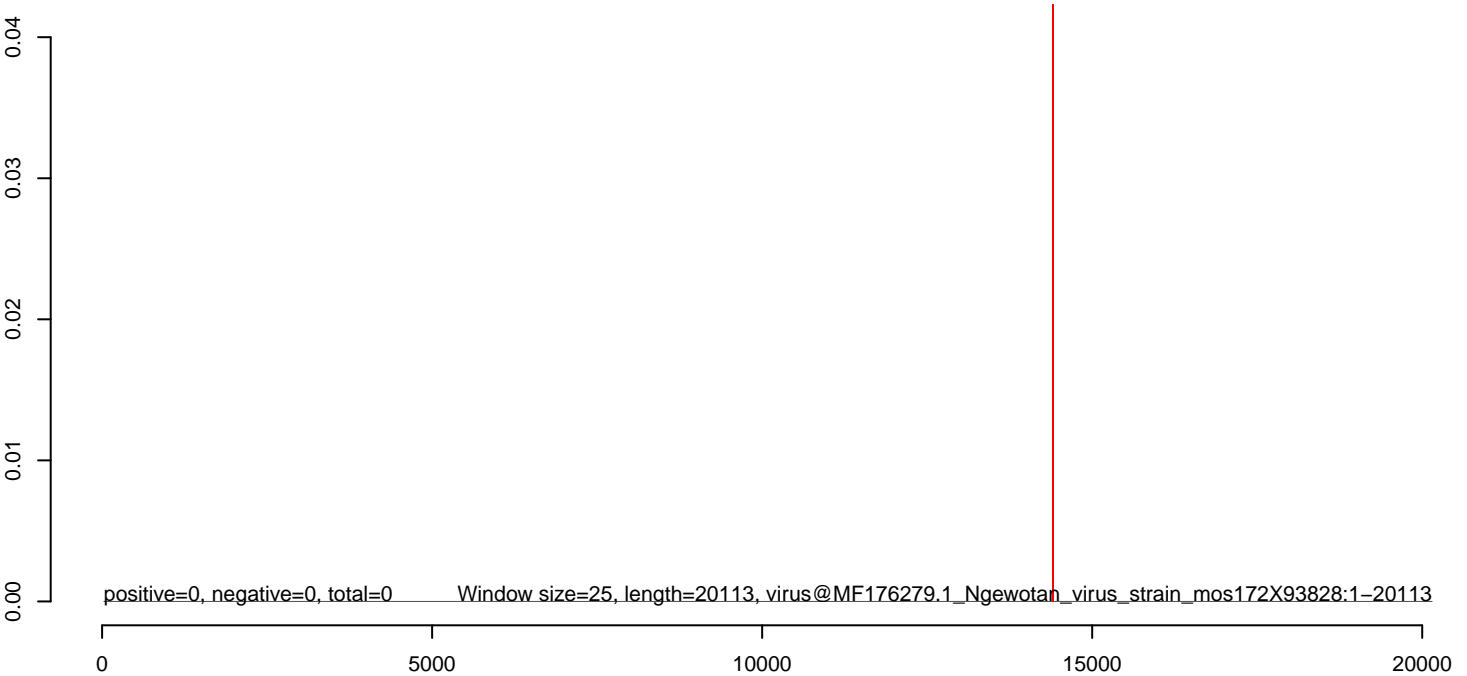
AeAlbo_C636_cells.18_23.rep



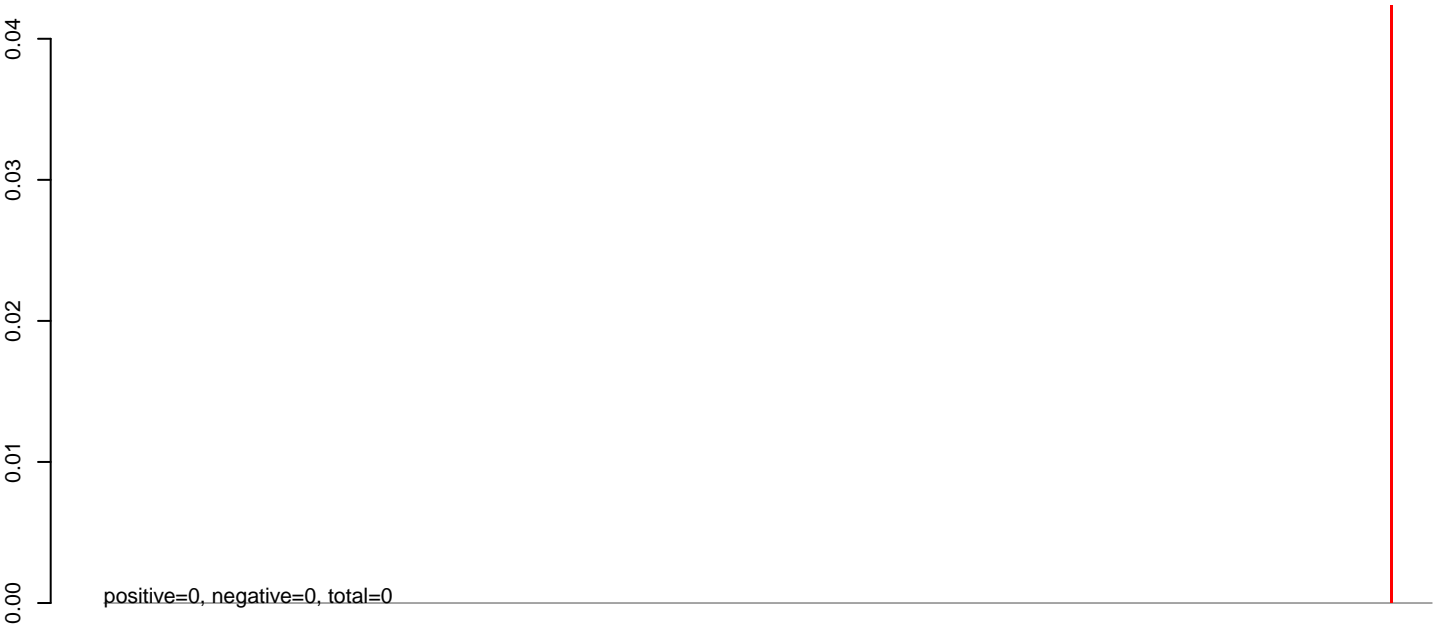
AeAlbo_C636_cells.24_35.rep



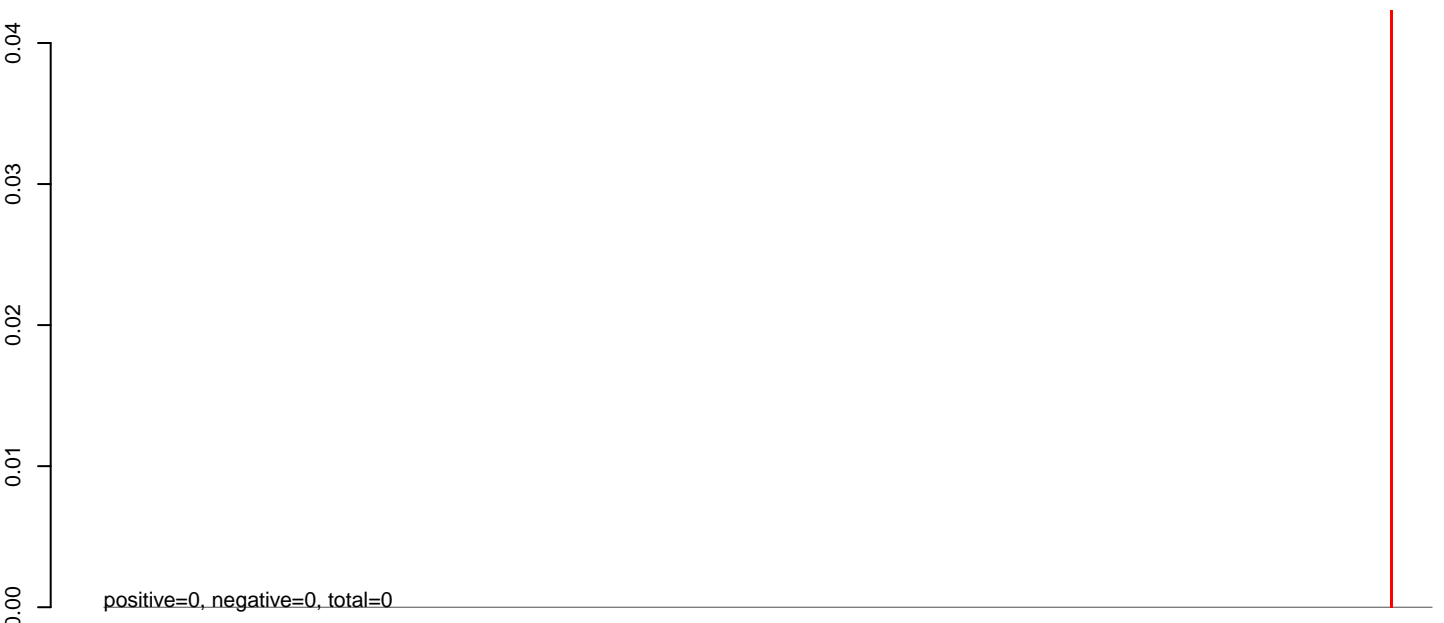
AeAlbo_C636_cells.rep



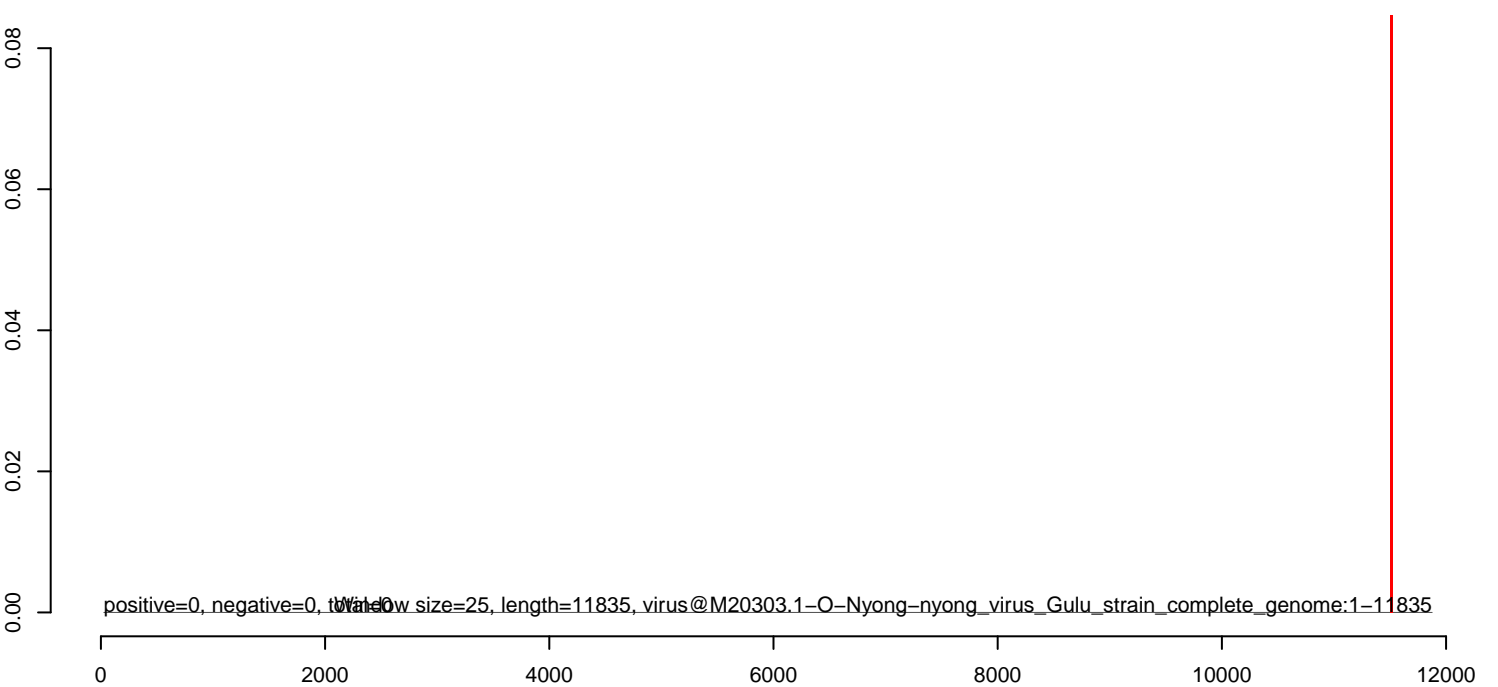
AeAlbo_C636_cells.18_23.rep



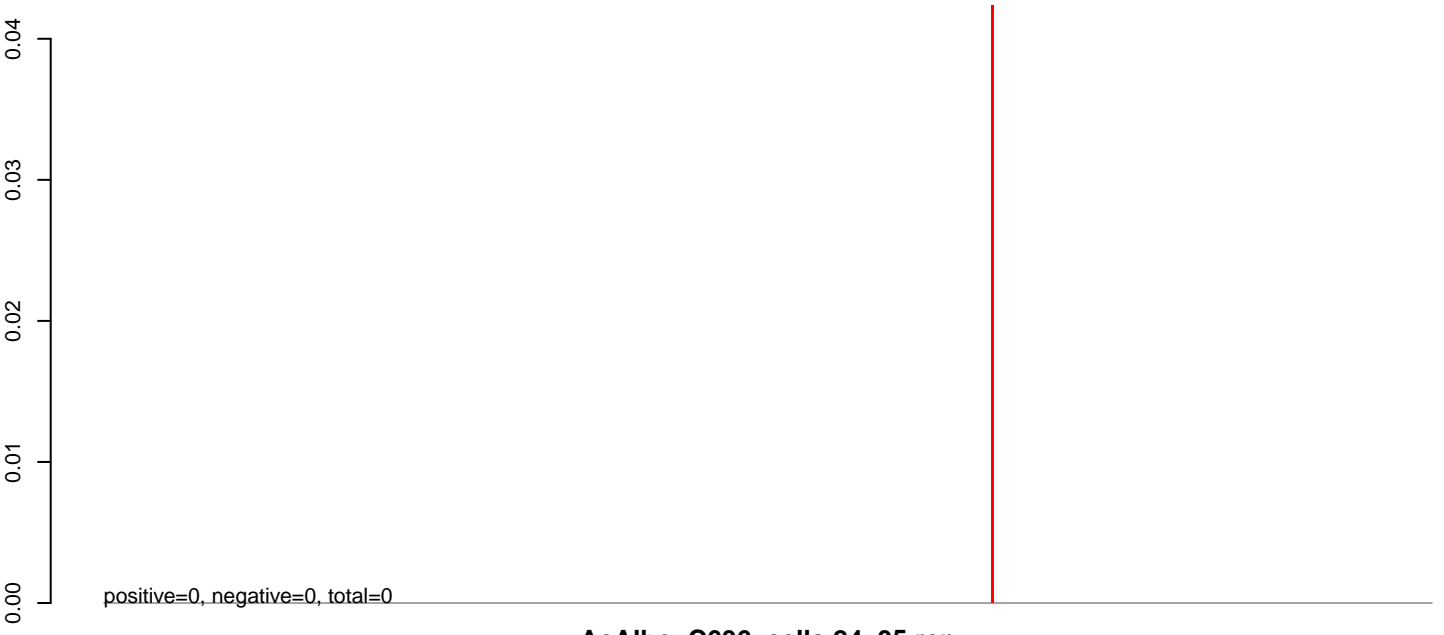
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



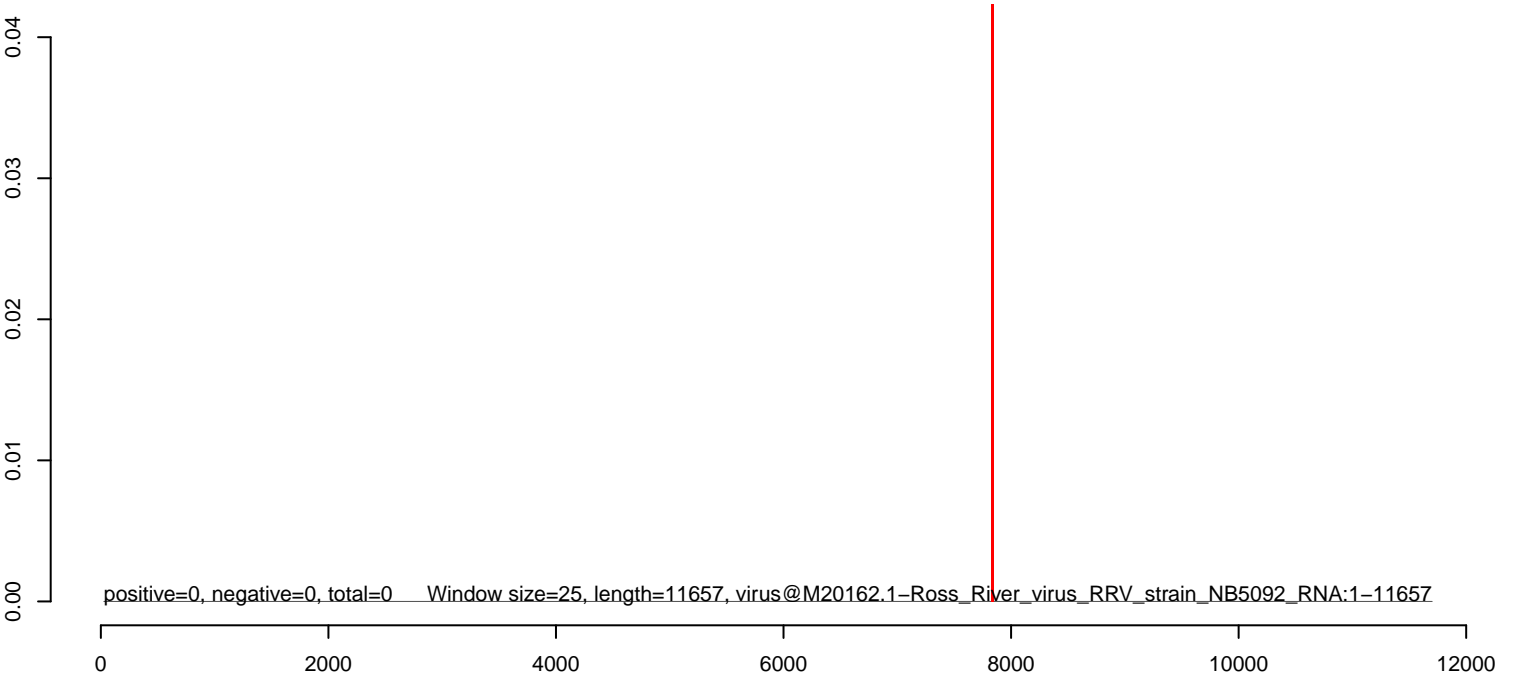
AeAlbo_C636_cells.18_23.rep



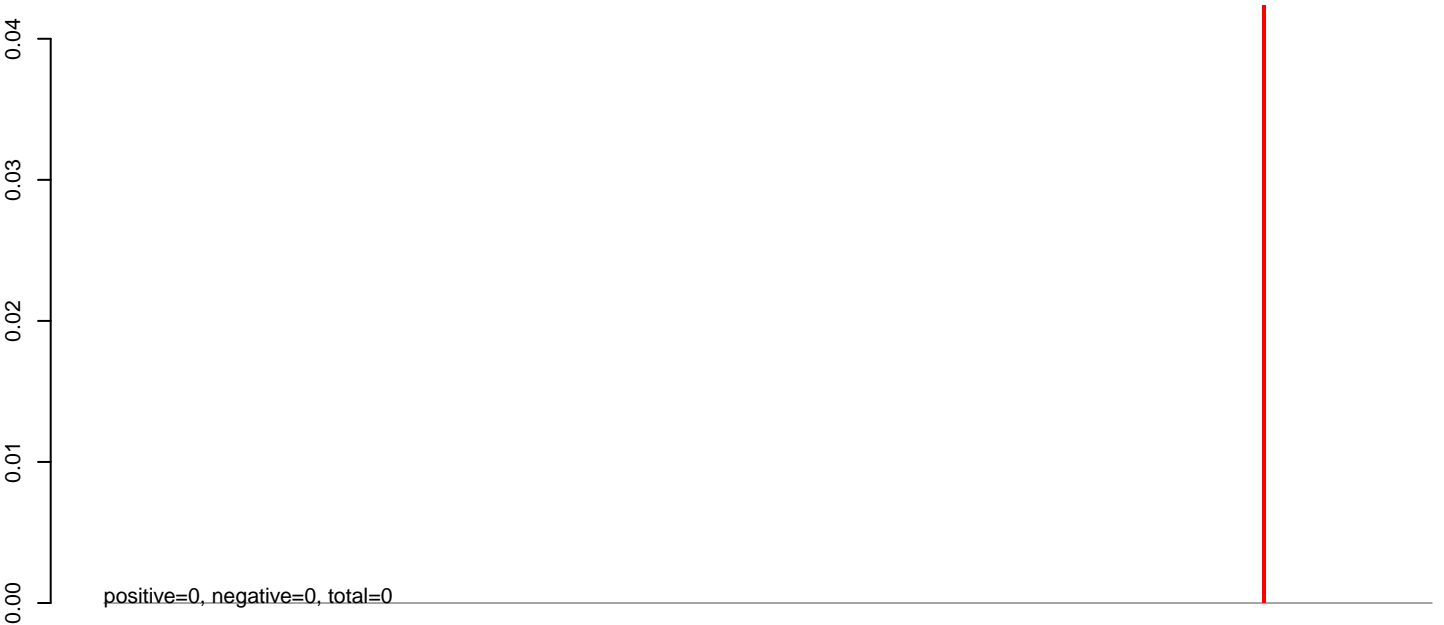
AeAlbo_C636_cells.24_35.rep



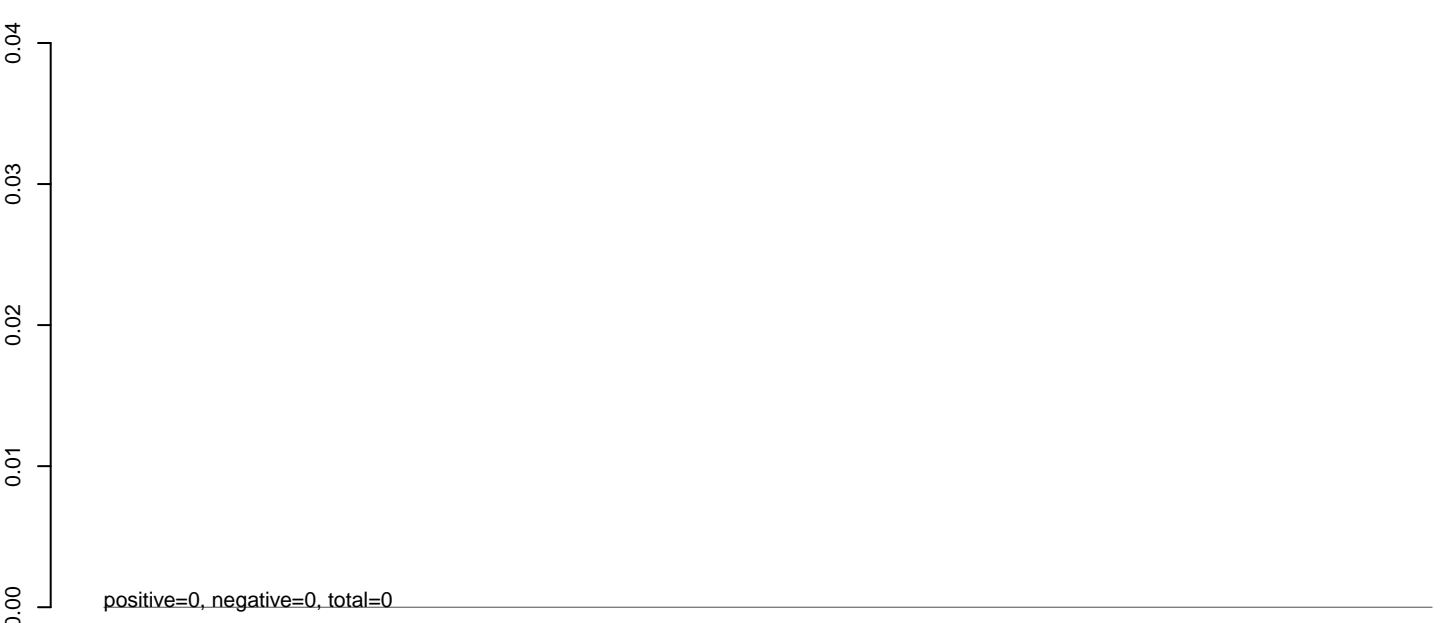
AeAlbo_C636_cells.rep



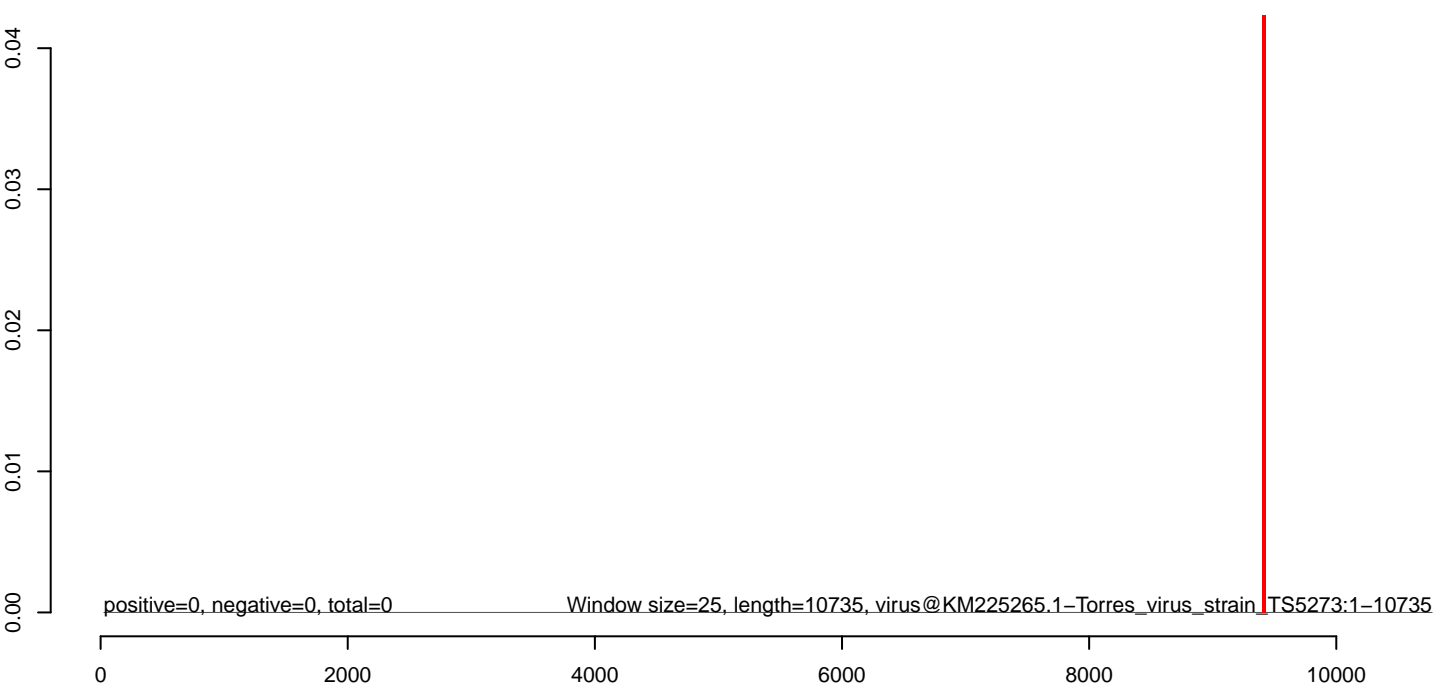
AeAlbo_C636_cells.18_23.rep



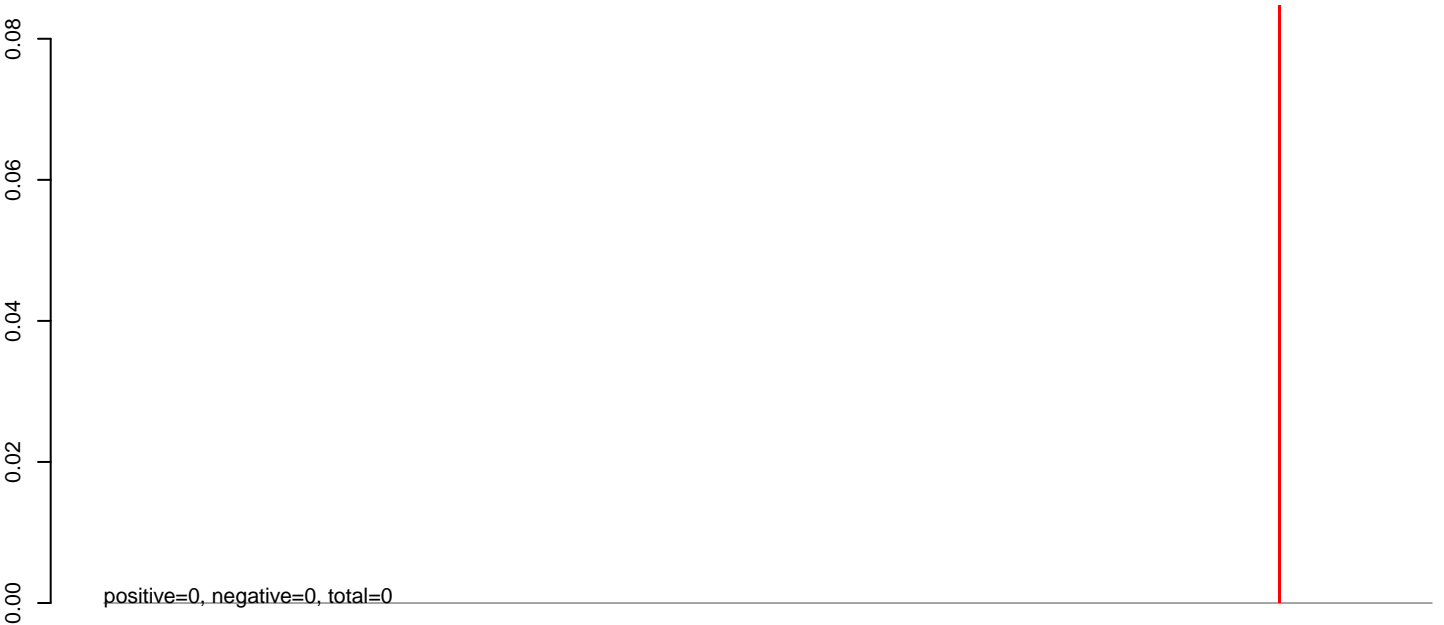
AeAlbo_C636_cells.24_35.rep



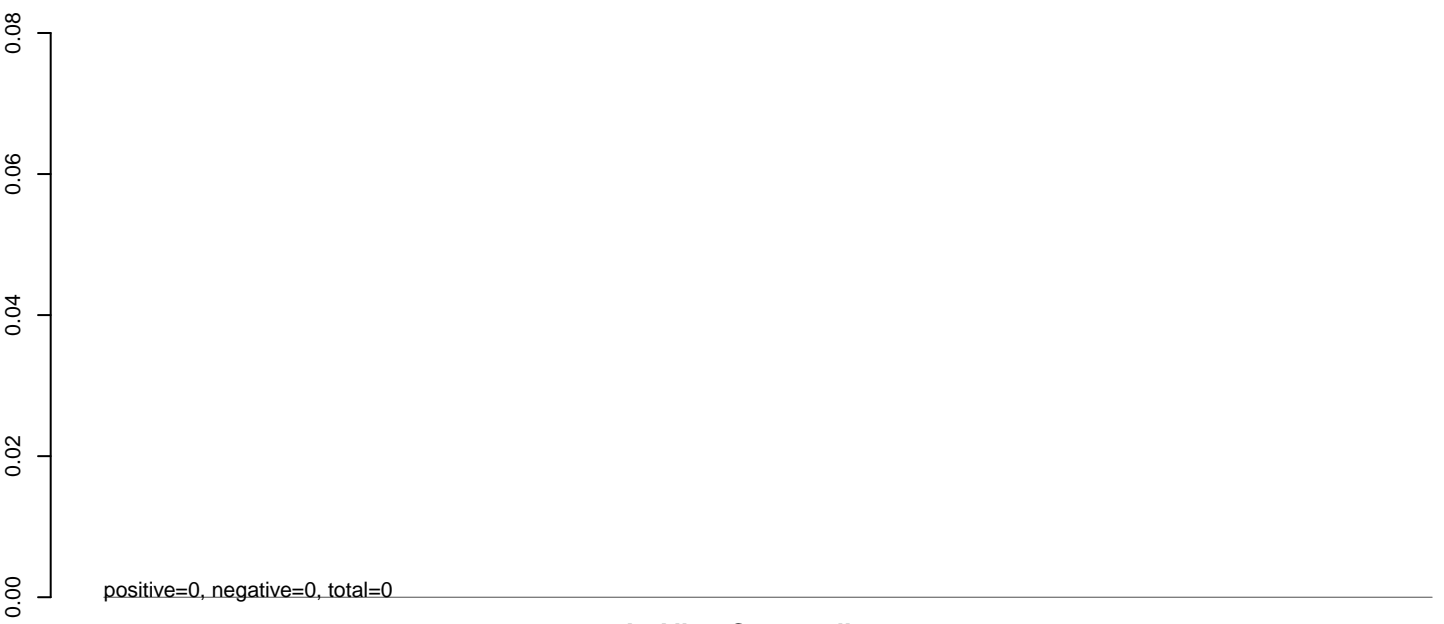
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



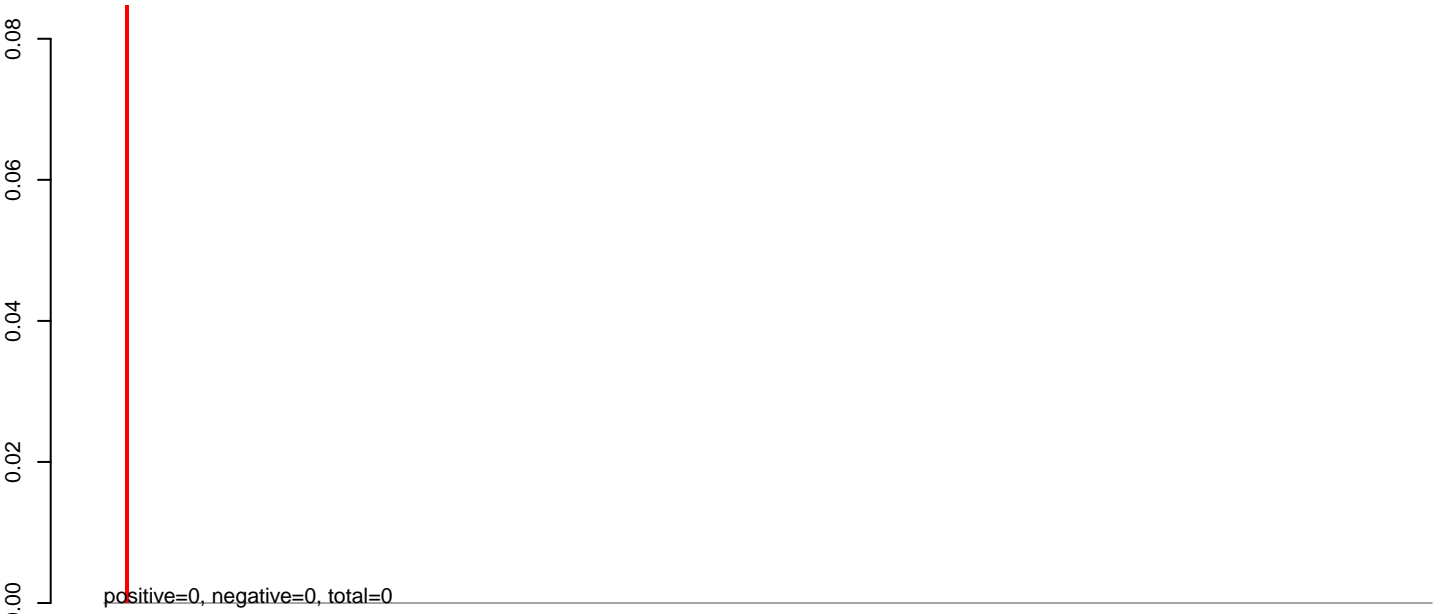
AeAlbo_C636_cells.24_35.rep



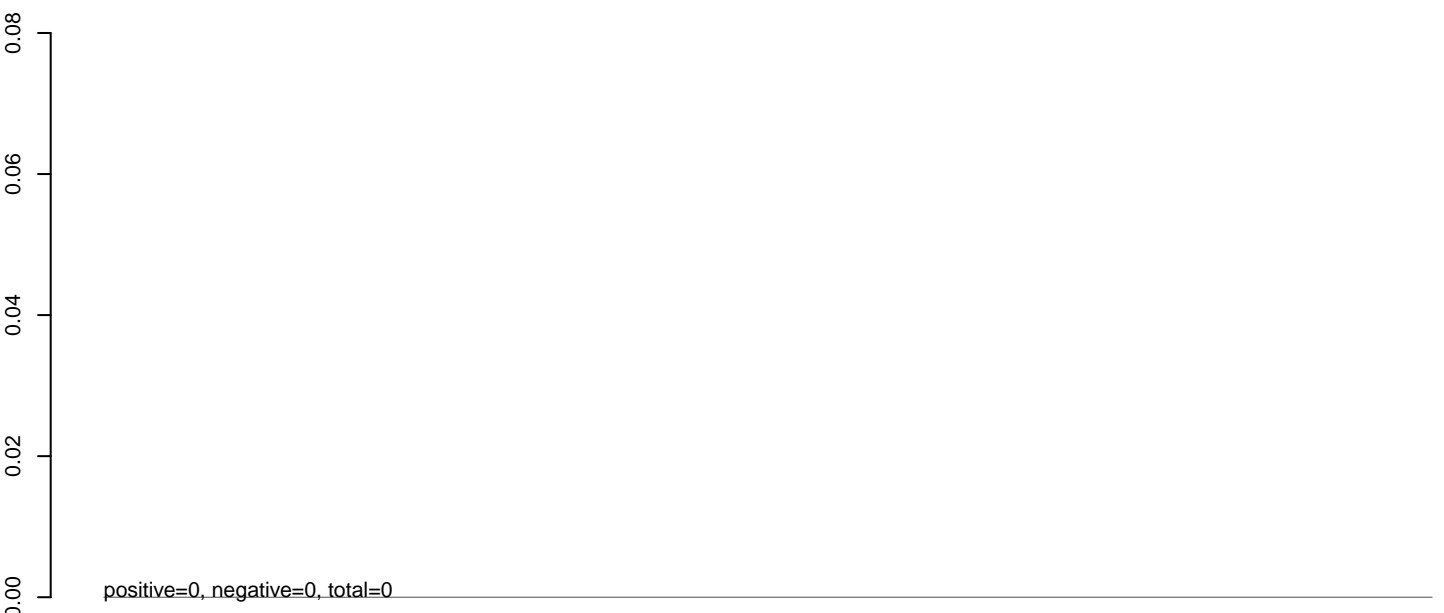
AeAlbo_C636_cells.rep



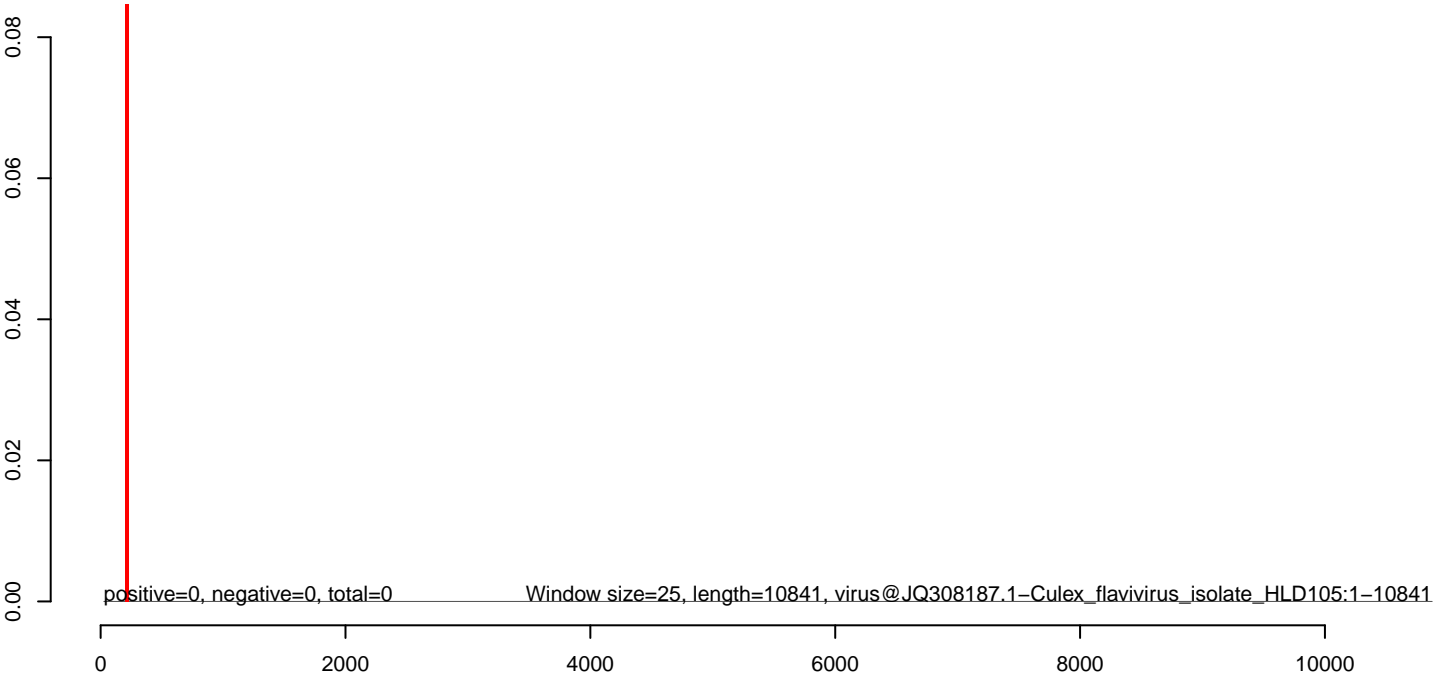
AeAlbo_C636_cells.18_23.rep



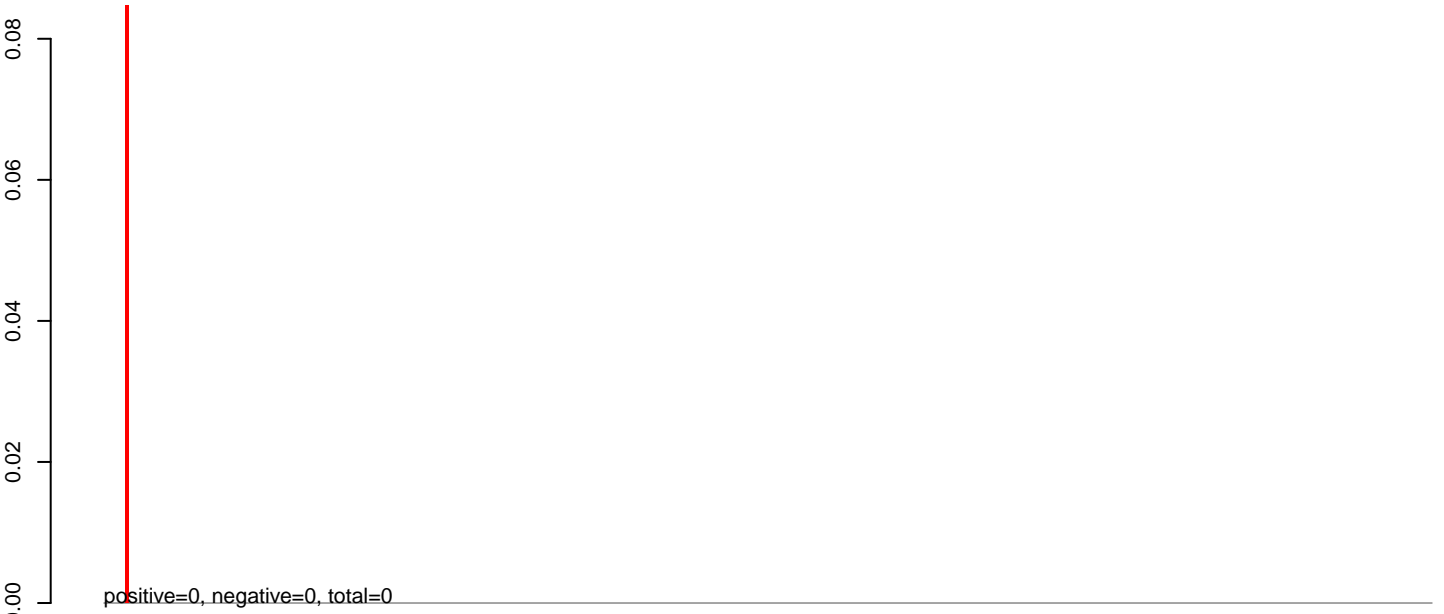
AeAlbo_C636_cells.24_35.rep



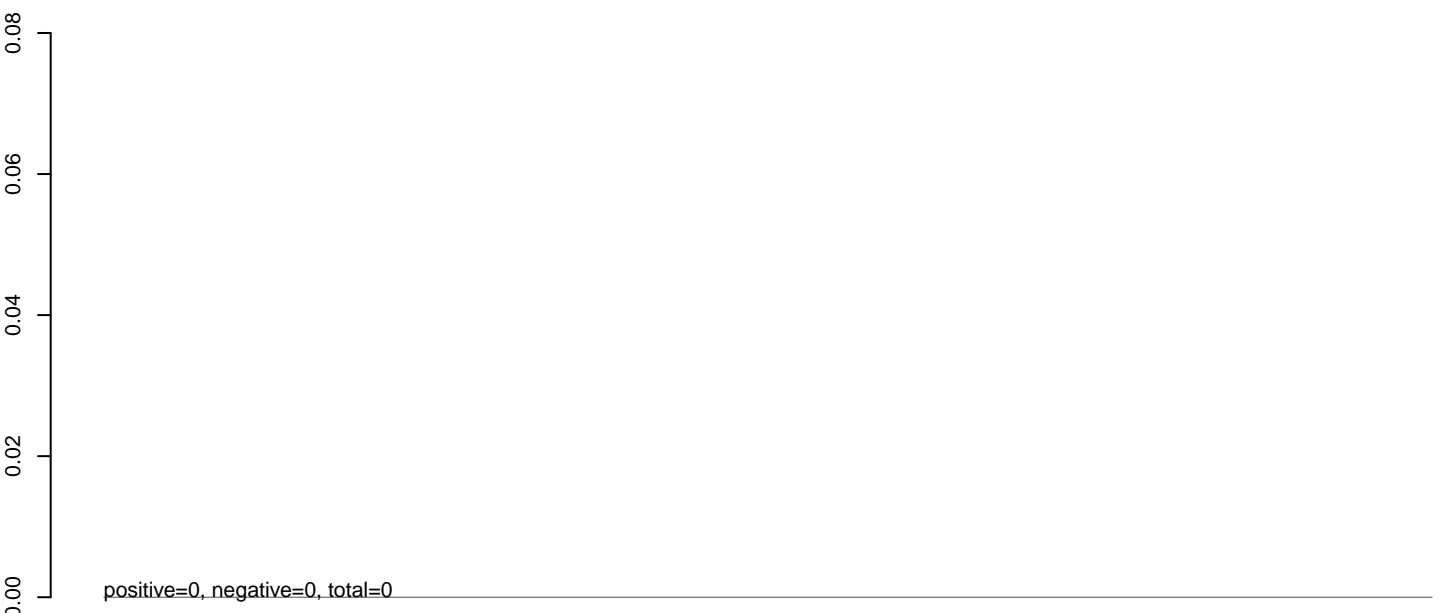
AeAlbo_C636_cells.rep



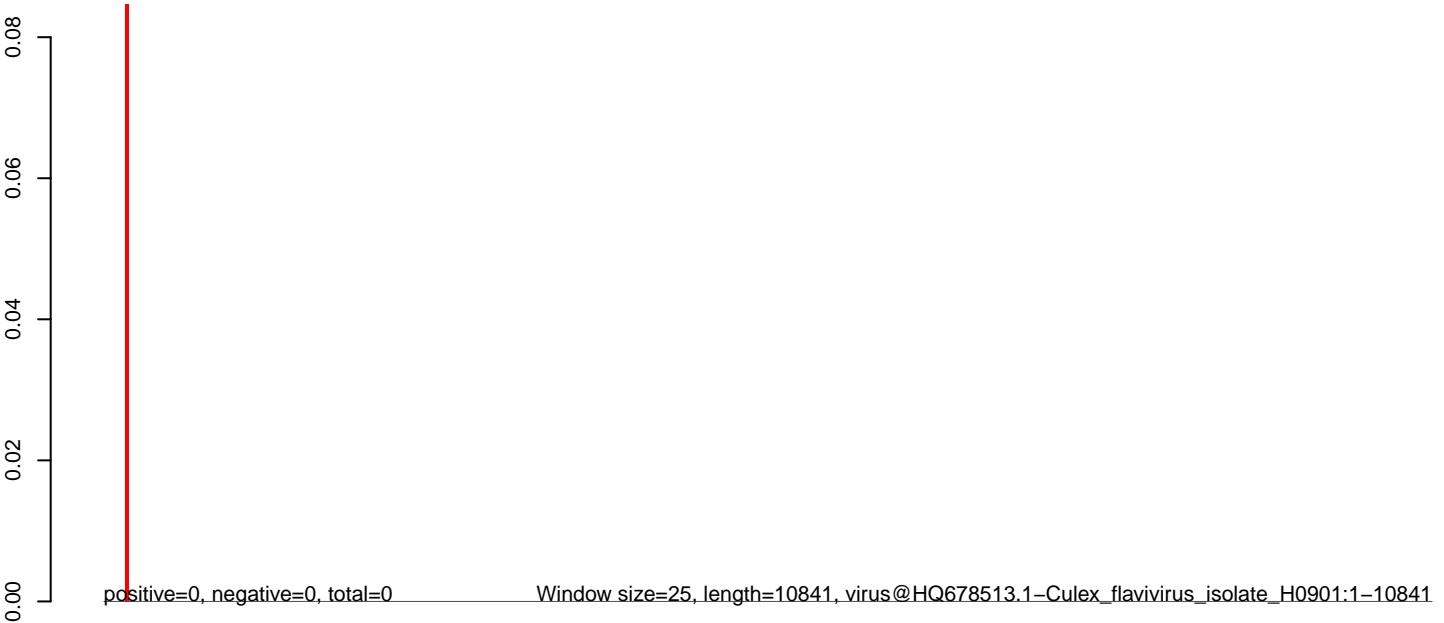
AeAlbo_C636_cells.18_23.rep



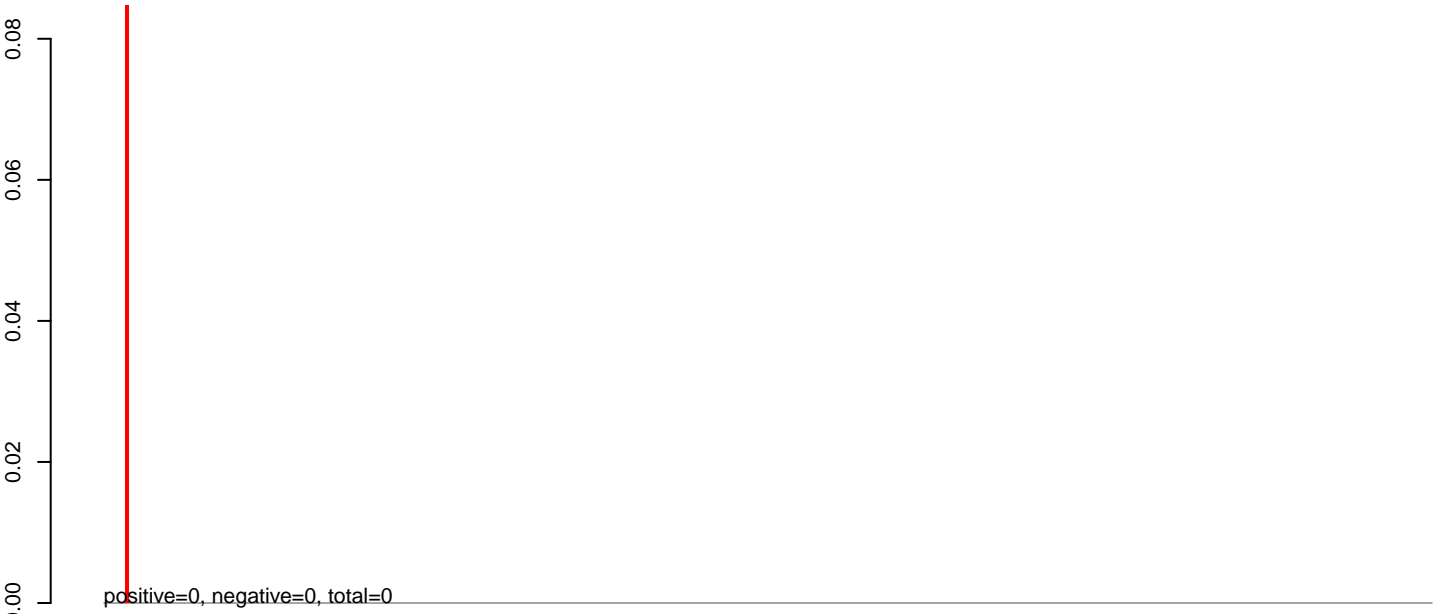
AeAlbo_C636_cells.24_35.rep



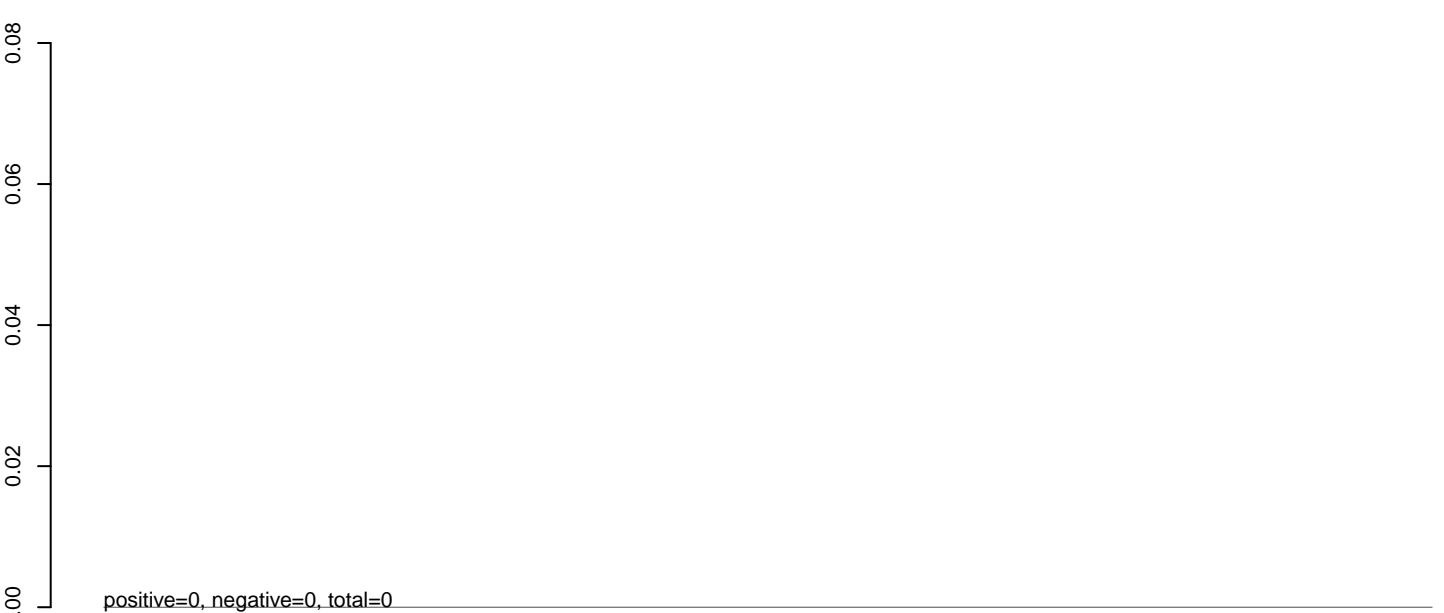
AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

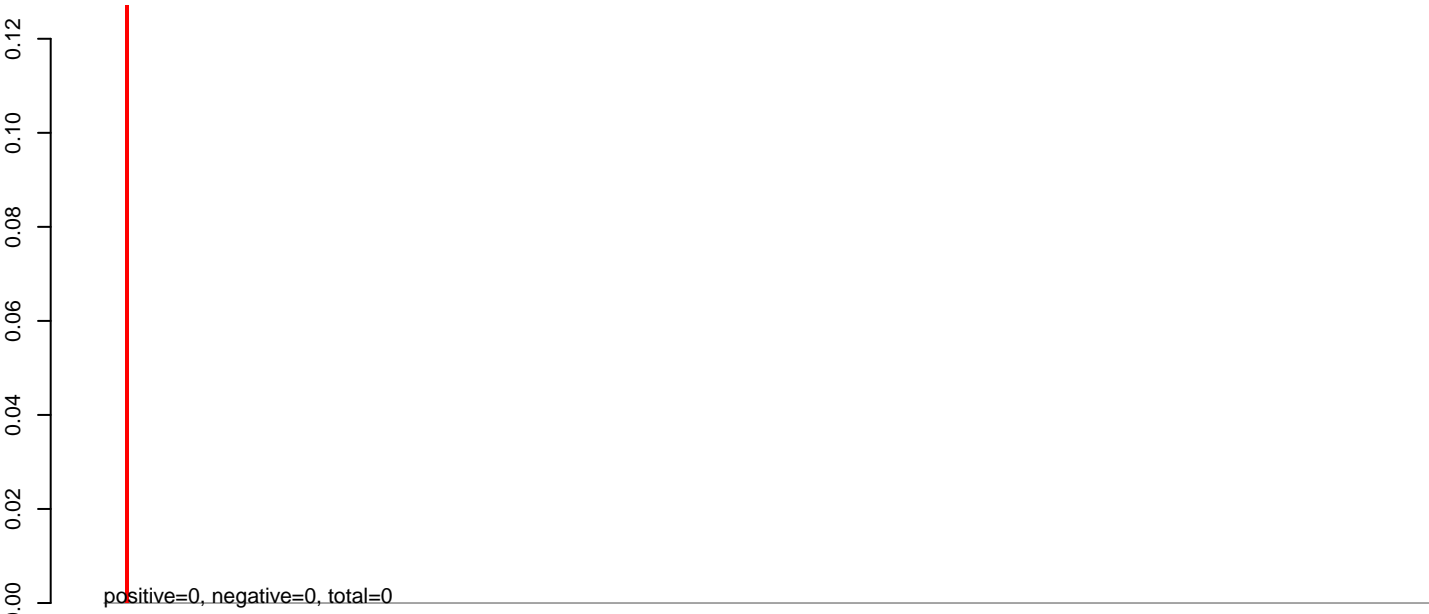


AeAlbo_C636_cells.rep

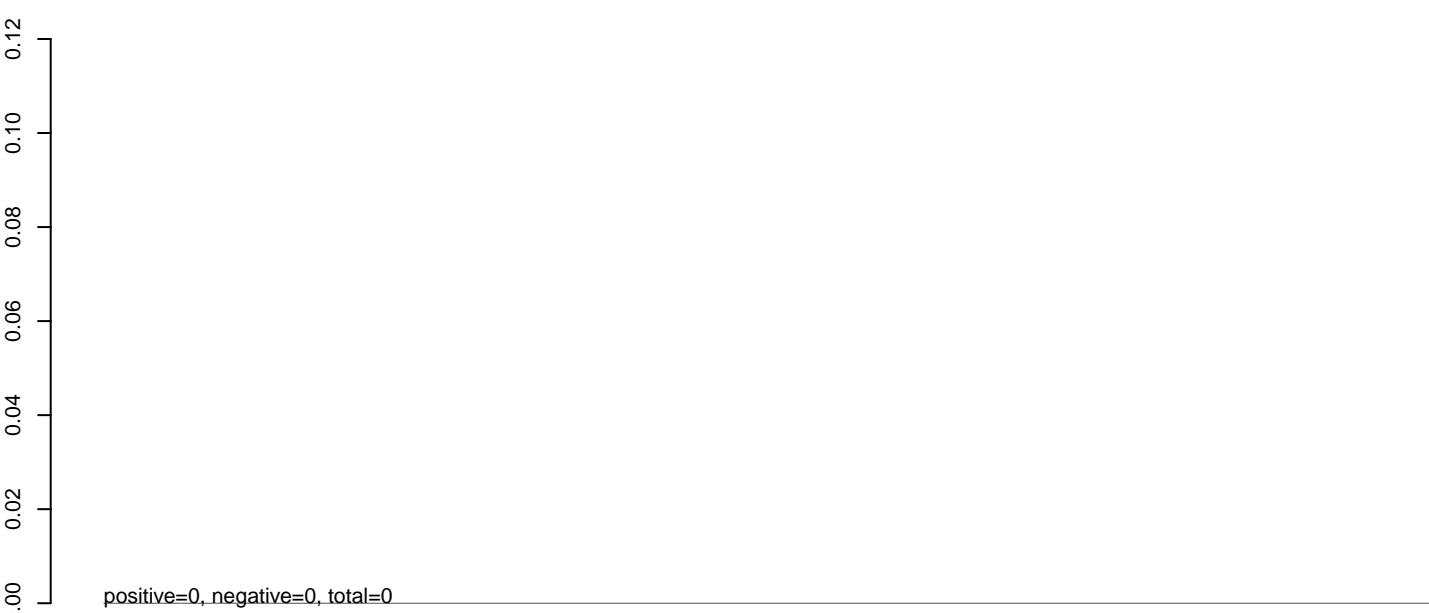


0 2000 4000 6000 8000 10000

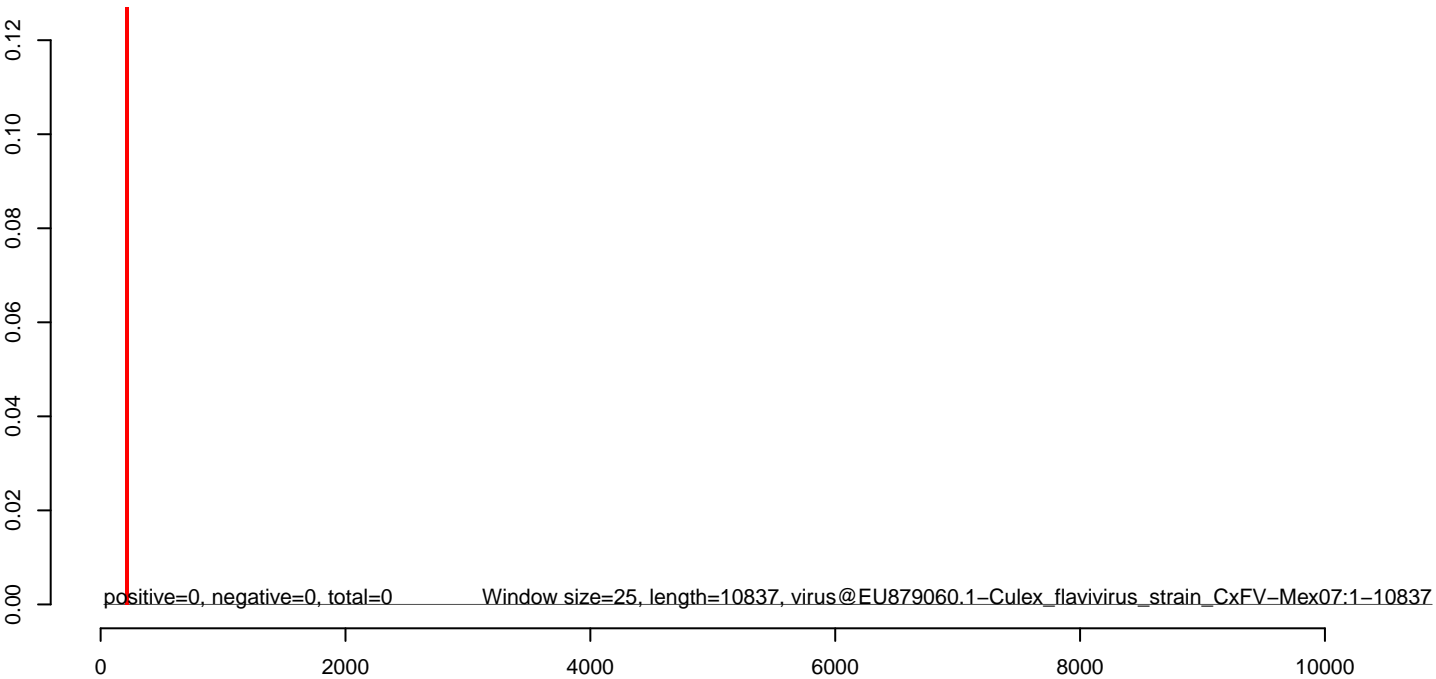
AeAlbo_C636_cells.18_23.rep



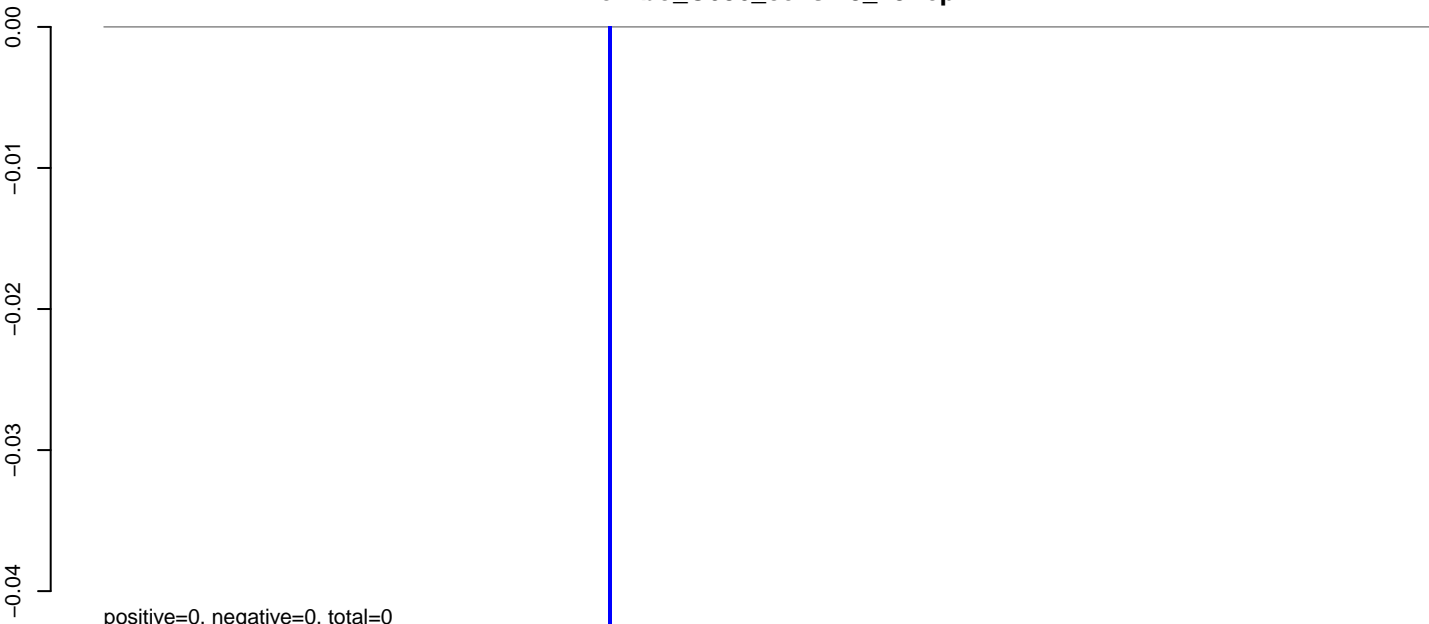
AeAlbo_C636_cells.24_35.rep



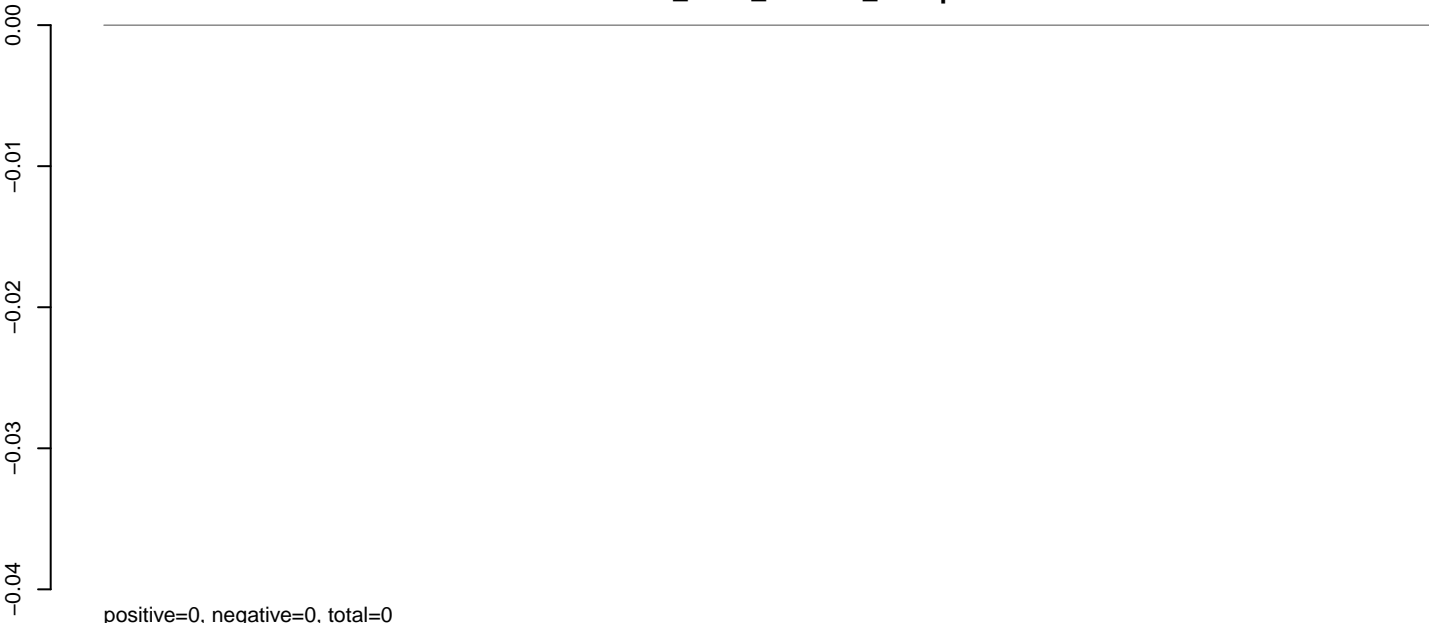
AeAlbo_C636_cells.rep



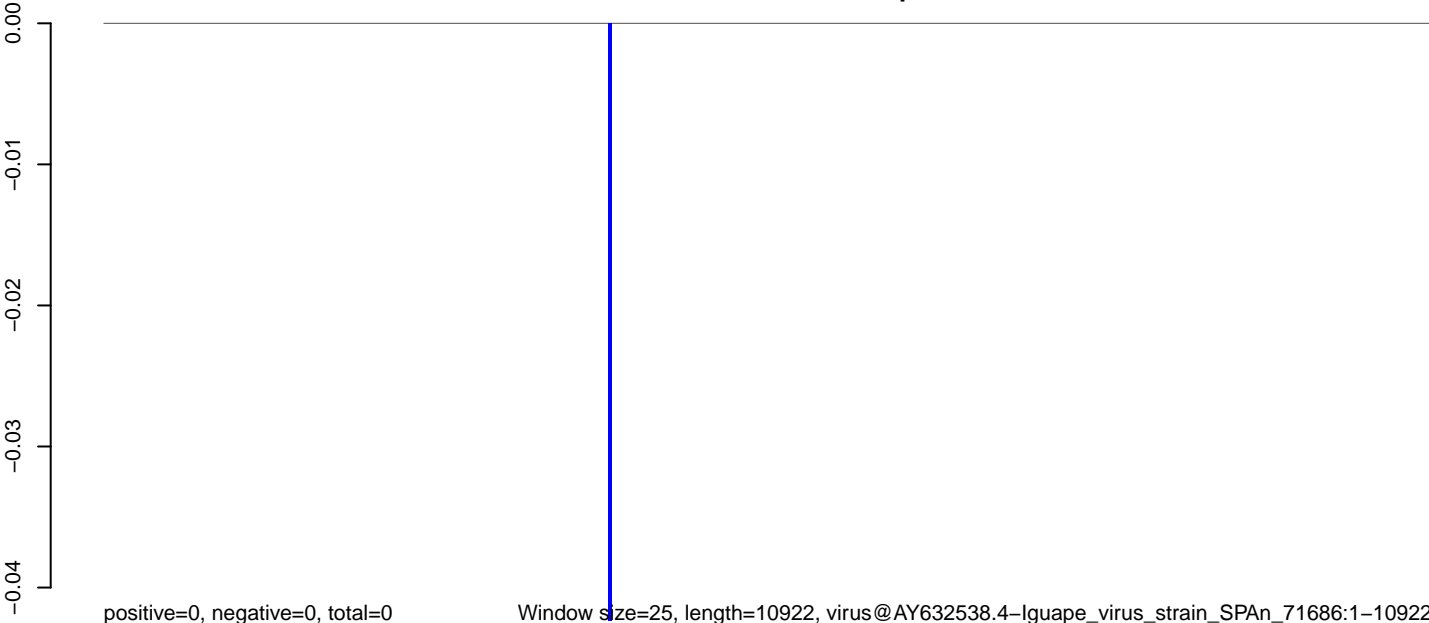
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

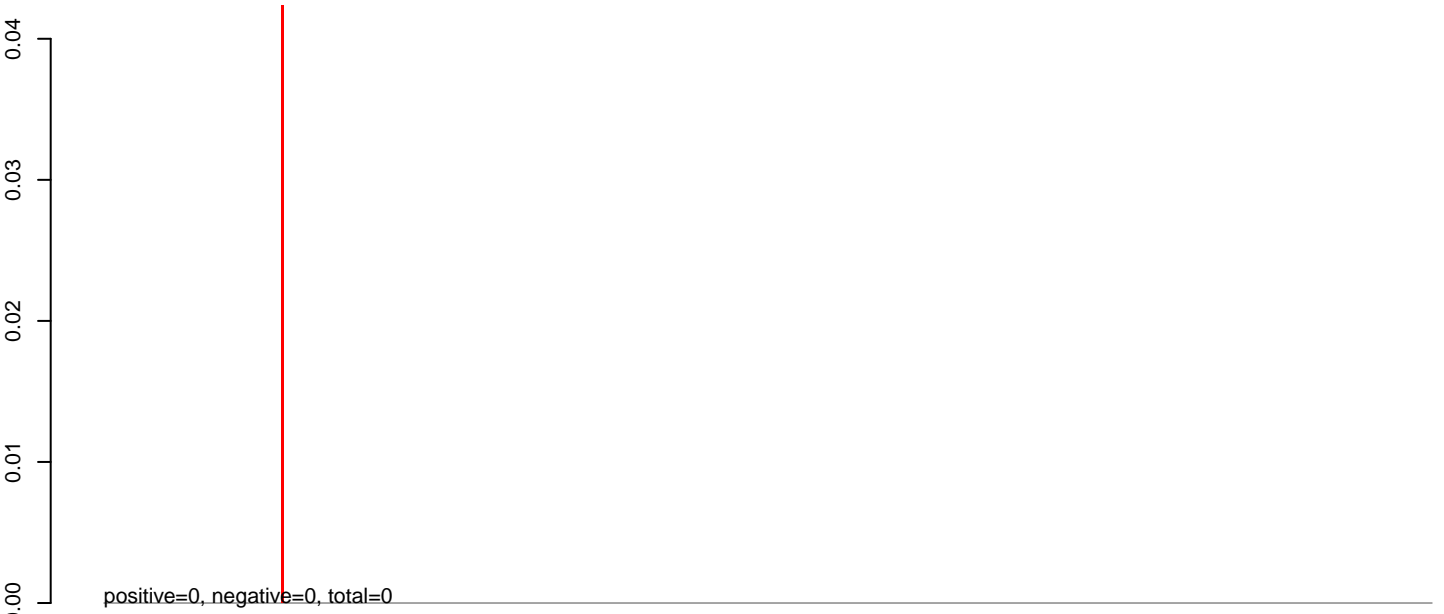


AeAlbo_C636_cells.rep

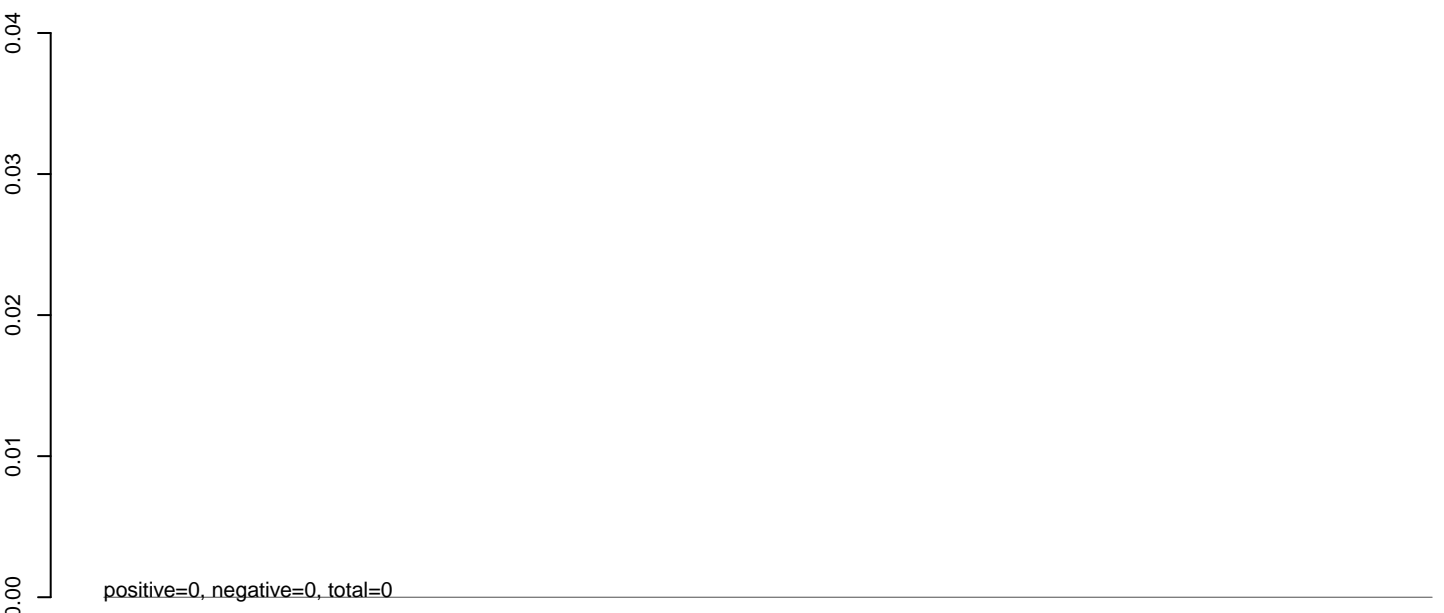


0 2000 4000 6000 8000 10000

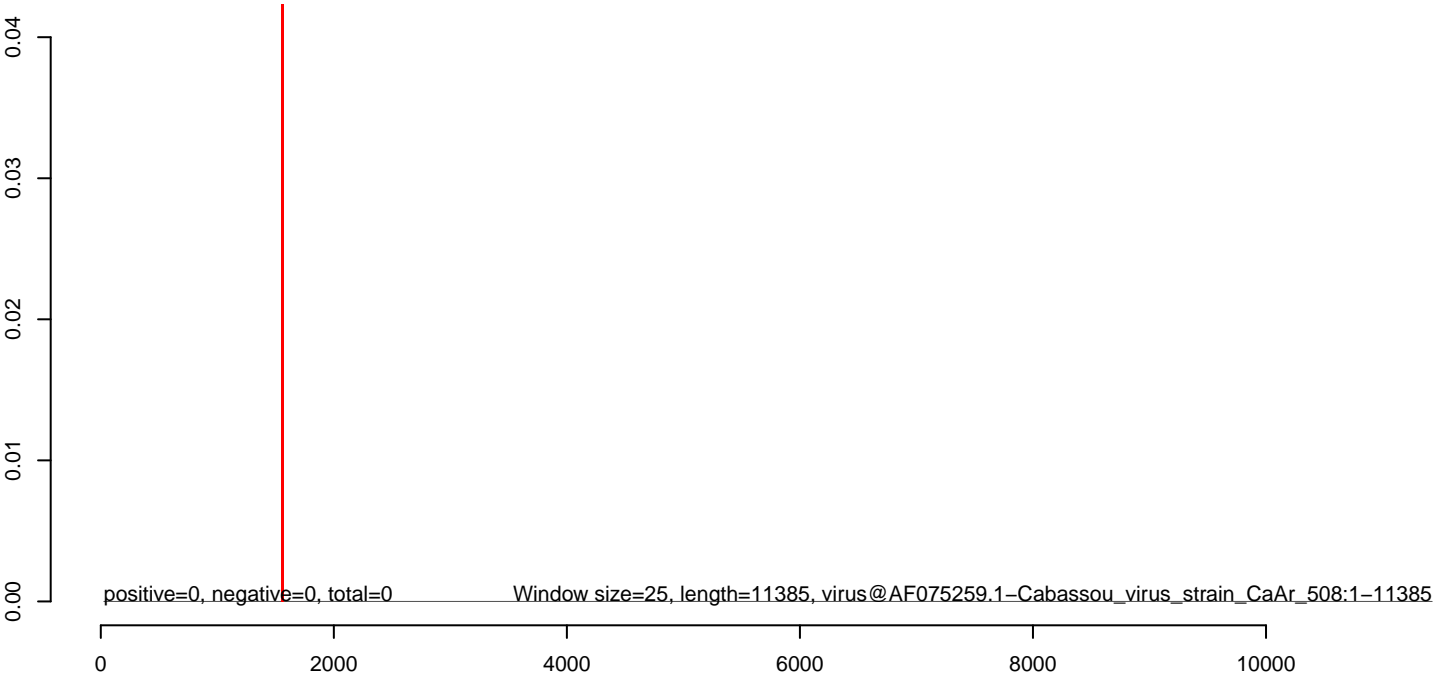
AeAlbo_C636_cells.18_23.rep



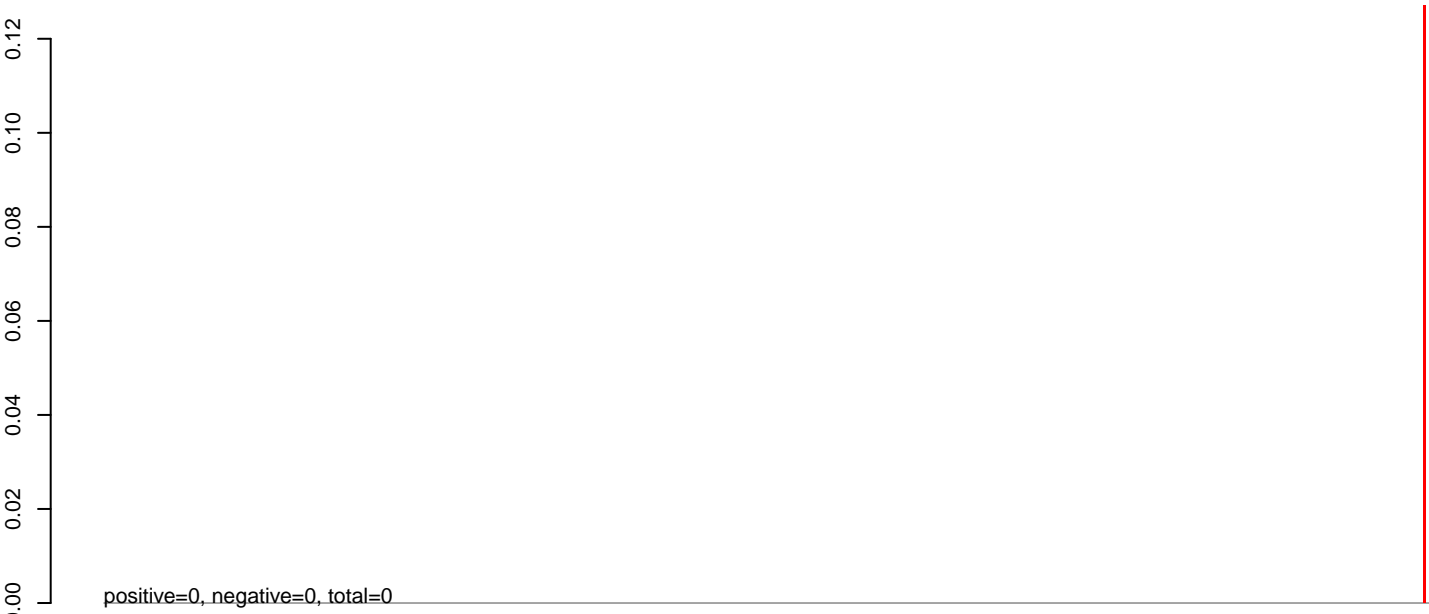
AeAlbo_C636_cells.24_35.rep



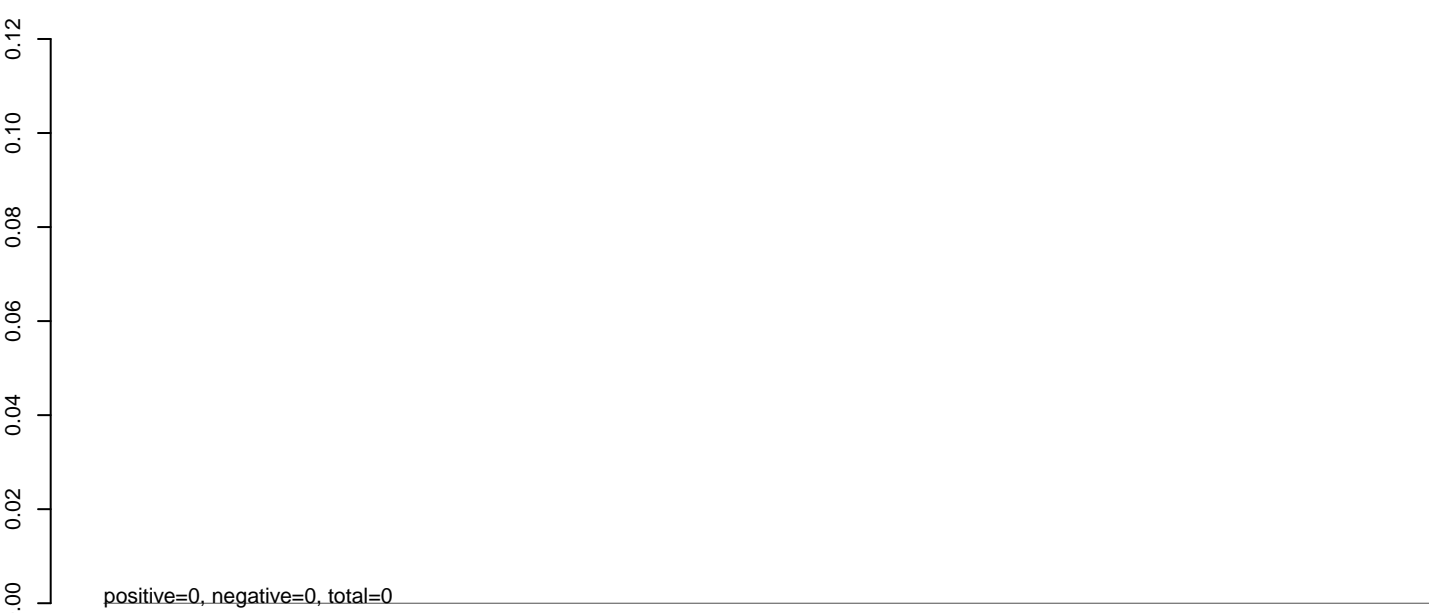
AeAlbo_C636_cells.rep



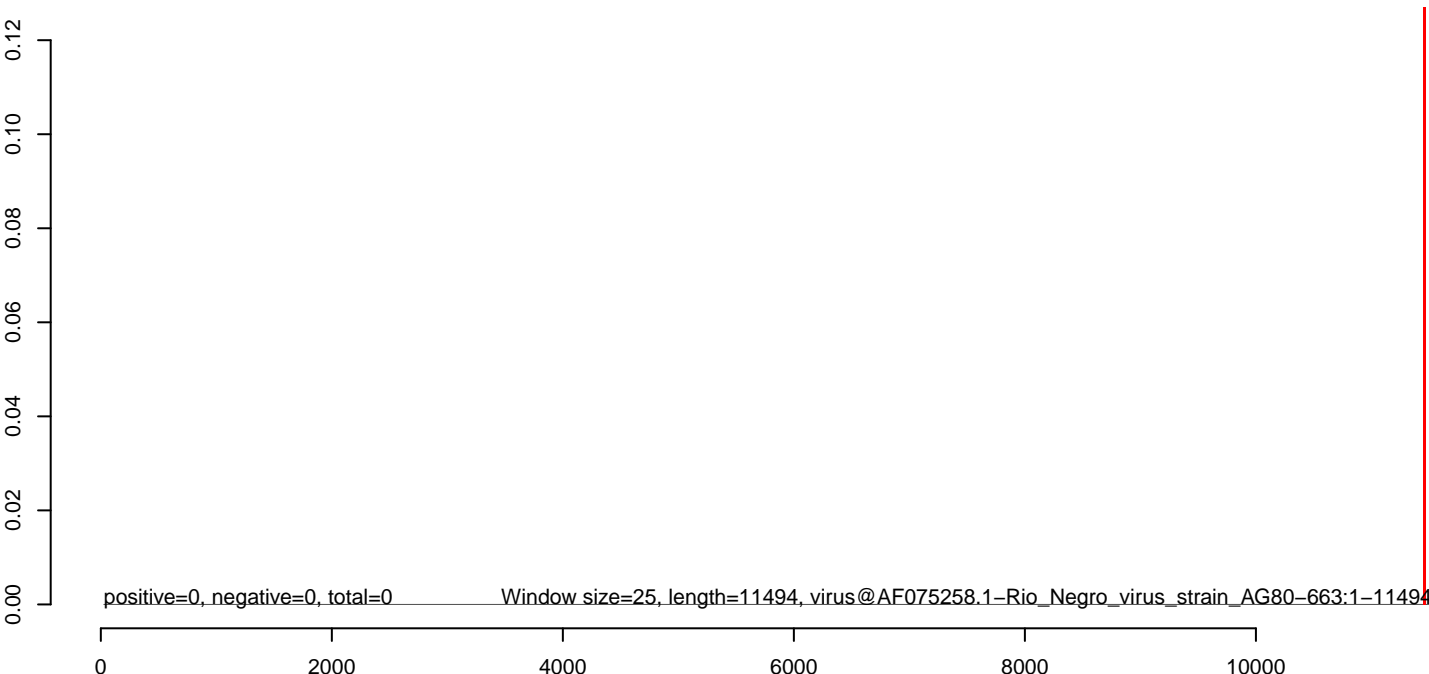
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



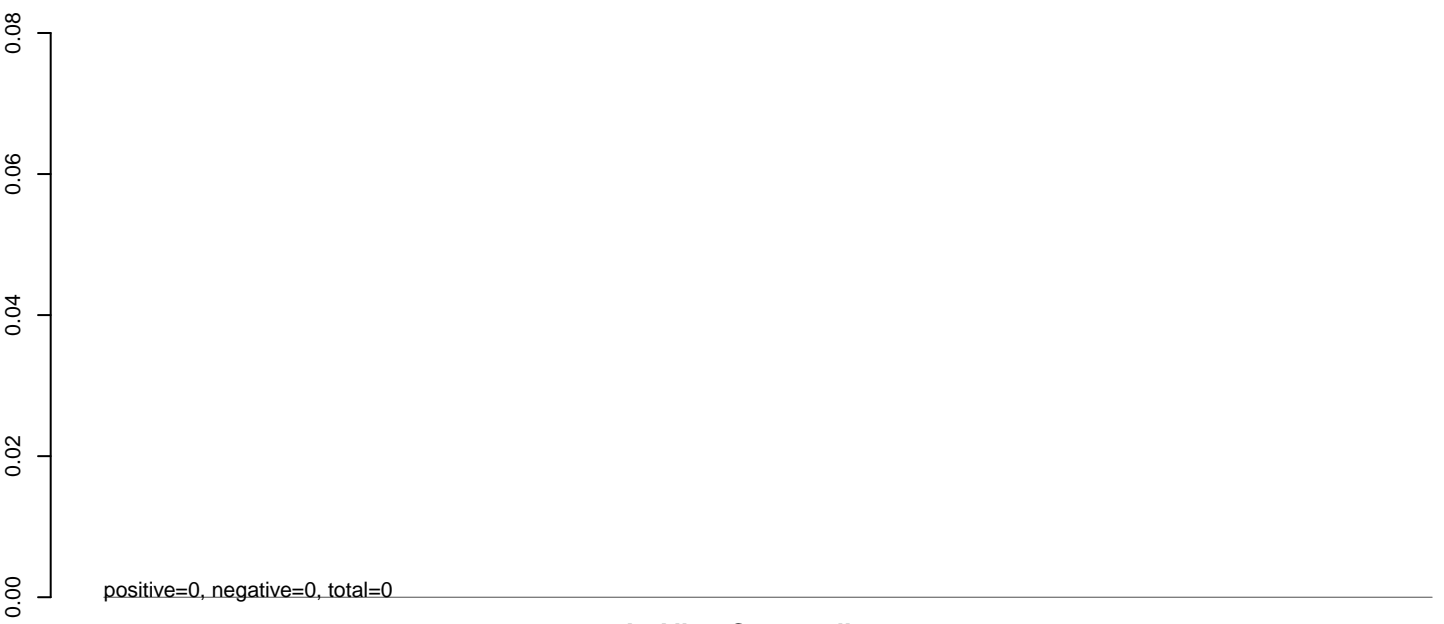
AeAlbo_C636_cells.rep



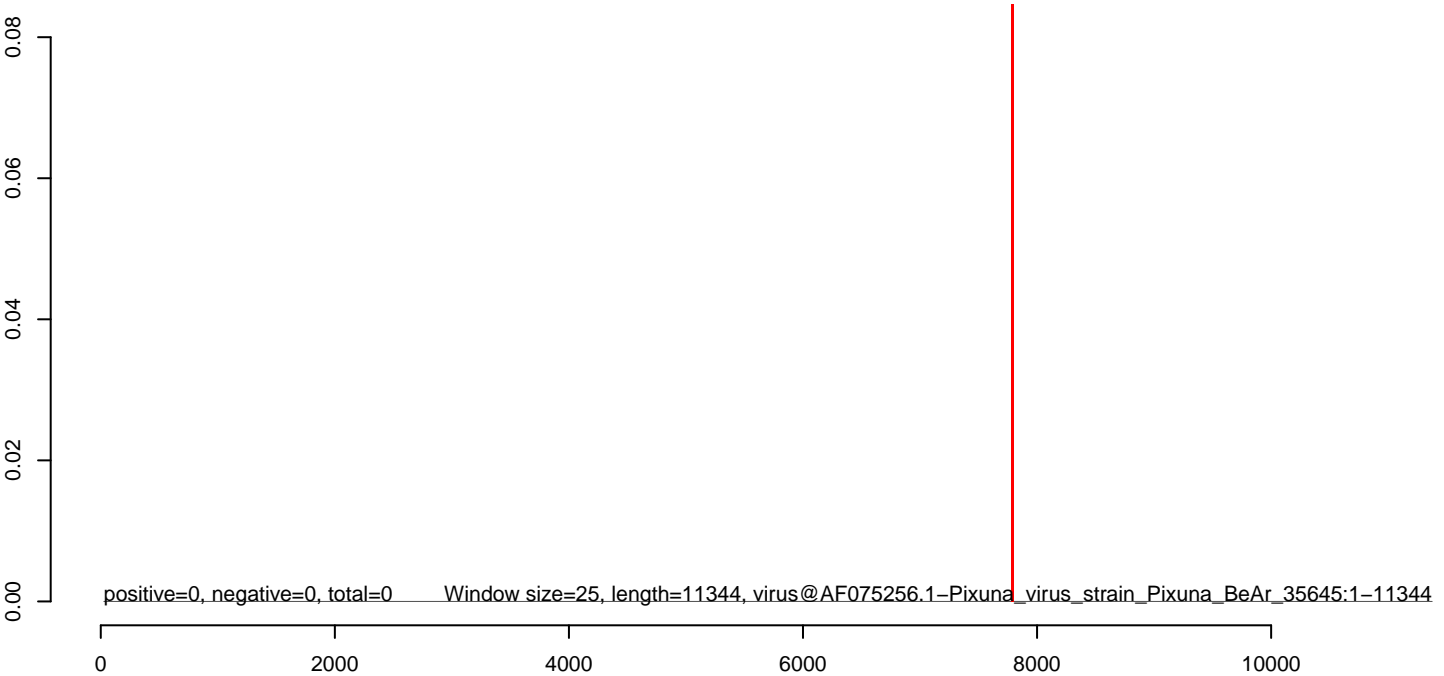
AeAlbo_C636_cells.18_23.rep



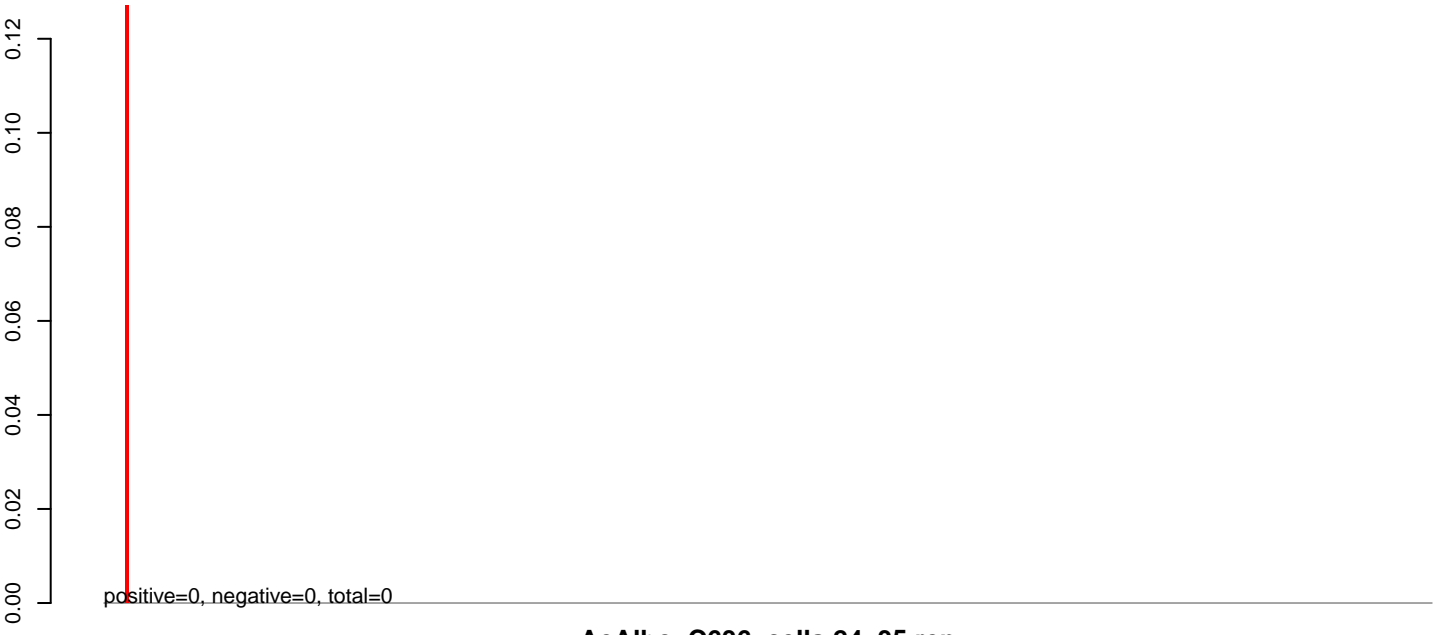
AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



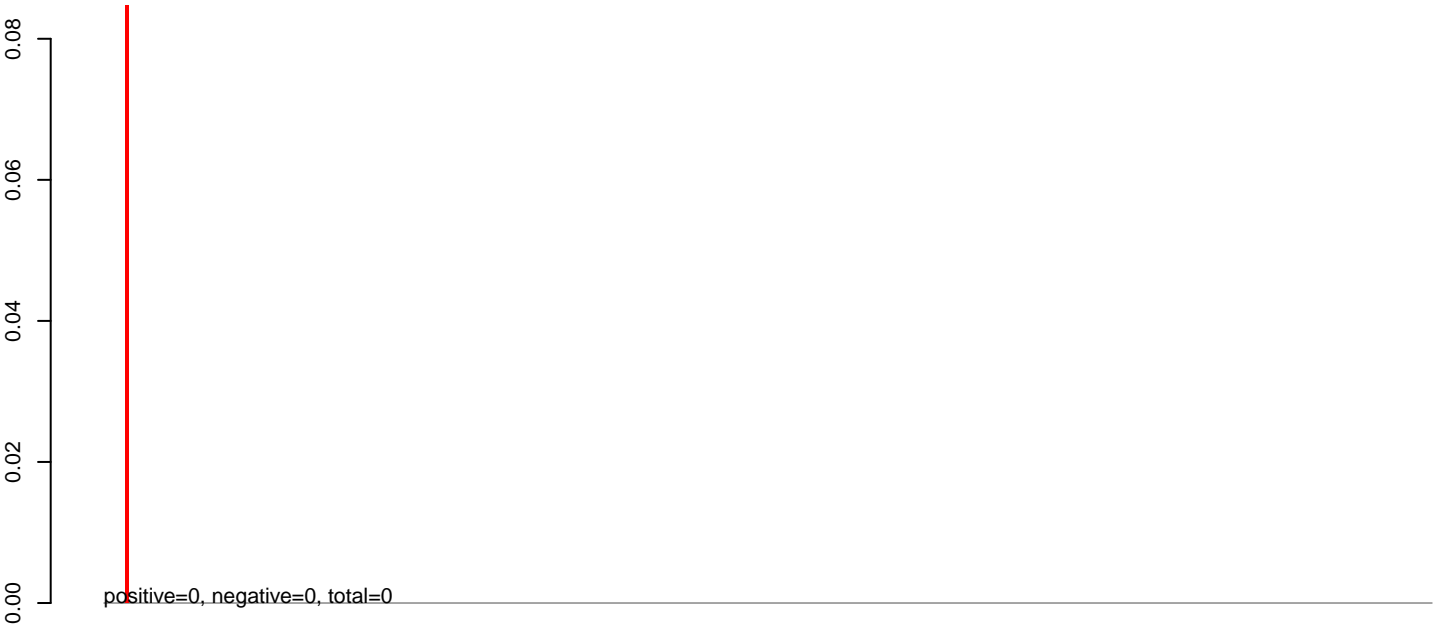
AeAlbo_C636_cells.24_35.rep



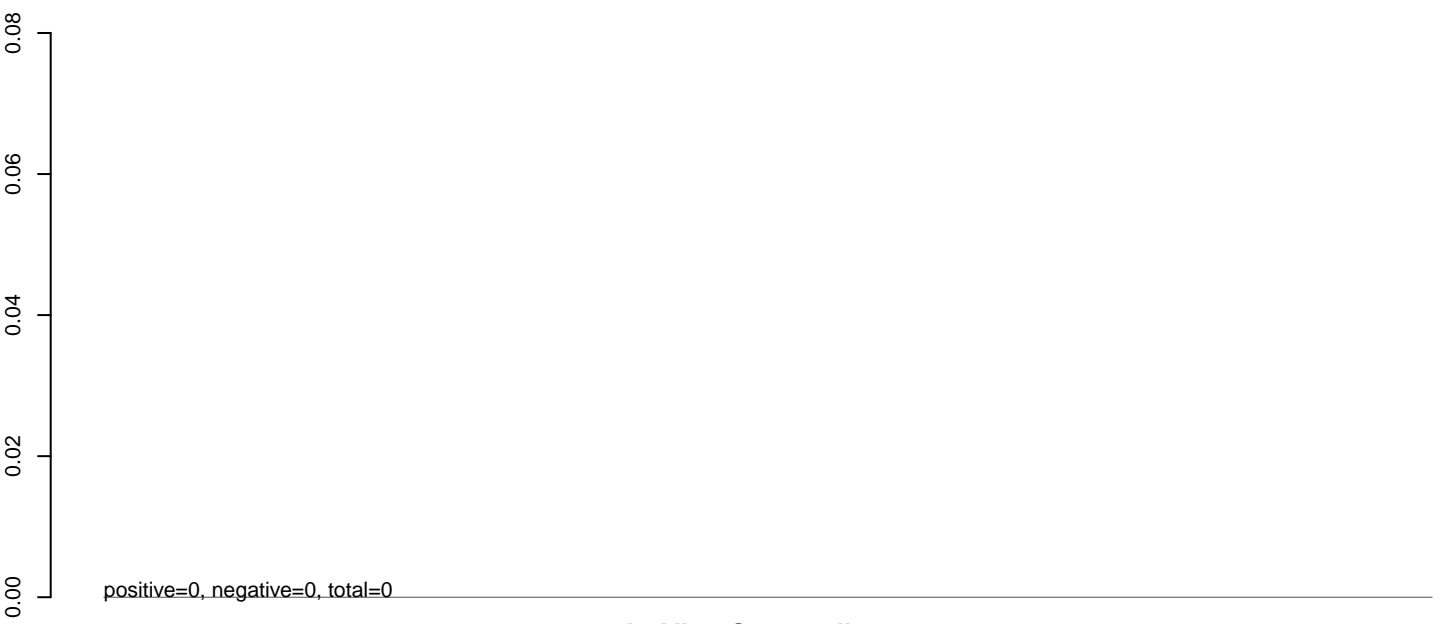
AeAlbo_C636_cells.rep



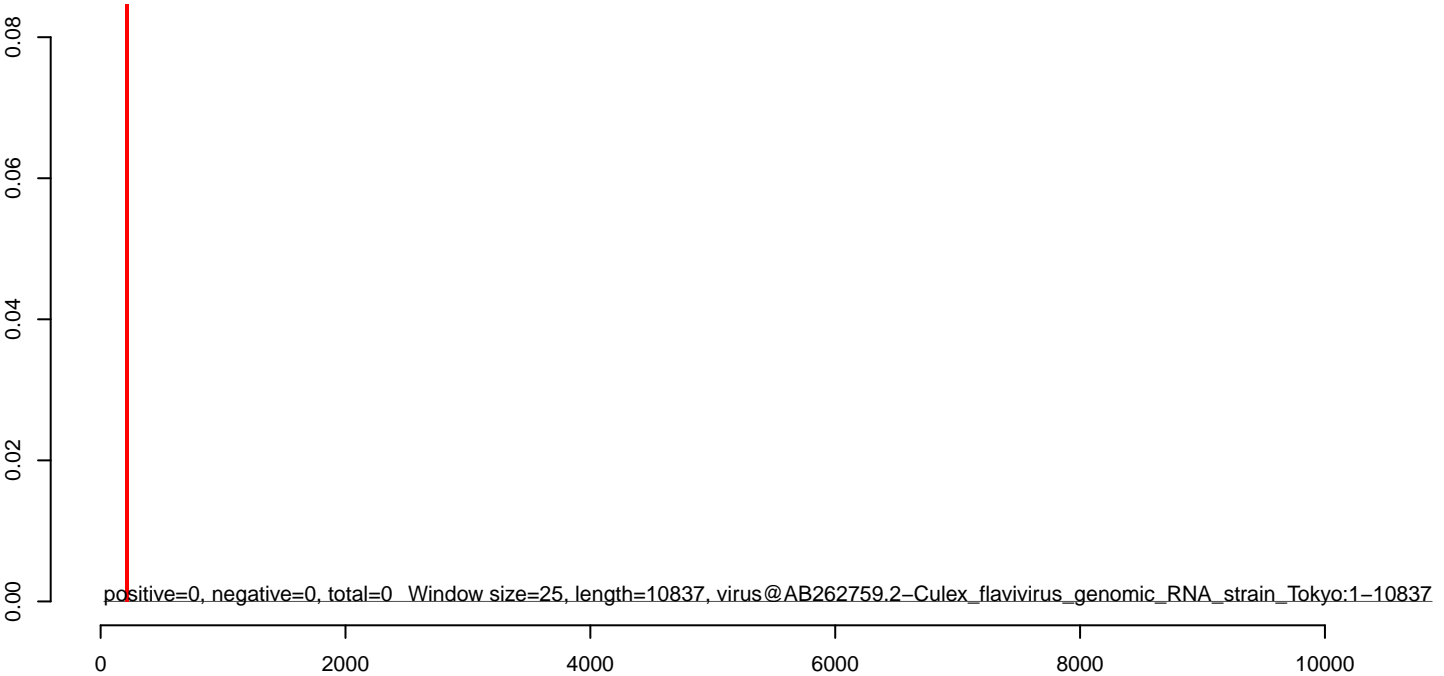
AeAlbo_C636_cells.18_23.rep



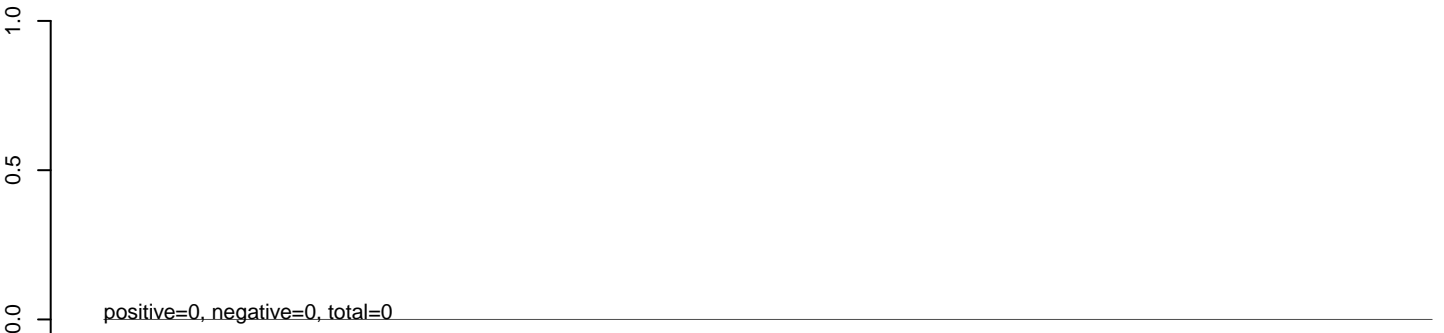
AeAlbo_C636_cells.24_35.rep



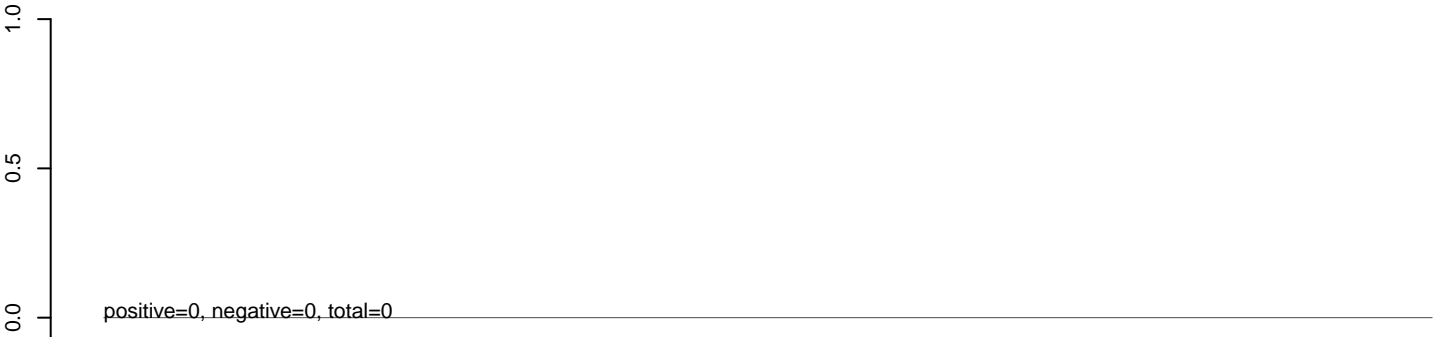
AeAlbo_C636_cells.rep



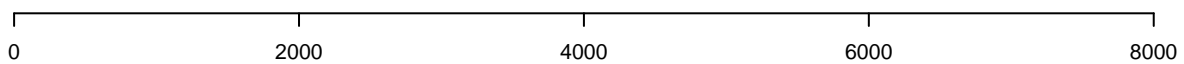
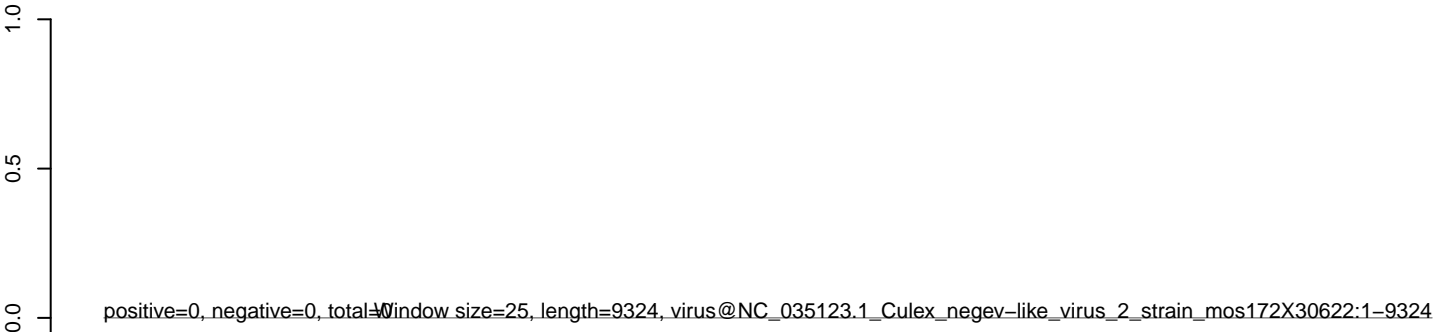
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



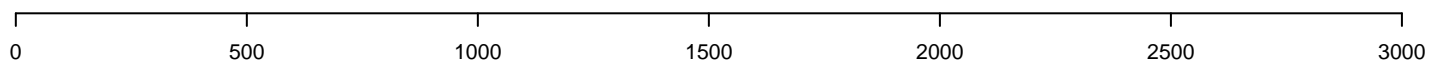
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



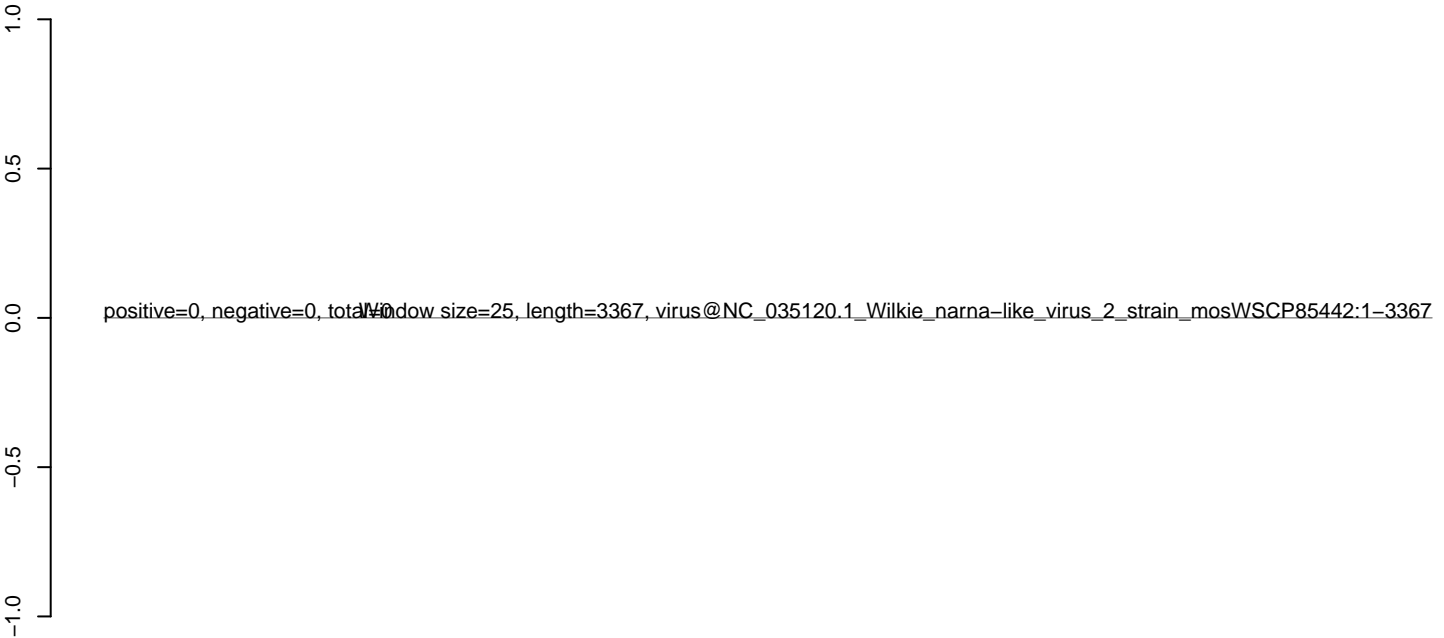
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

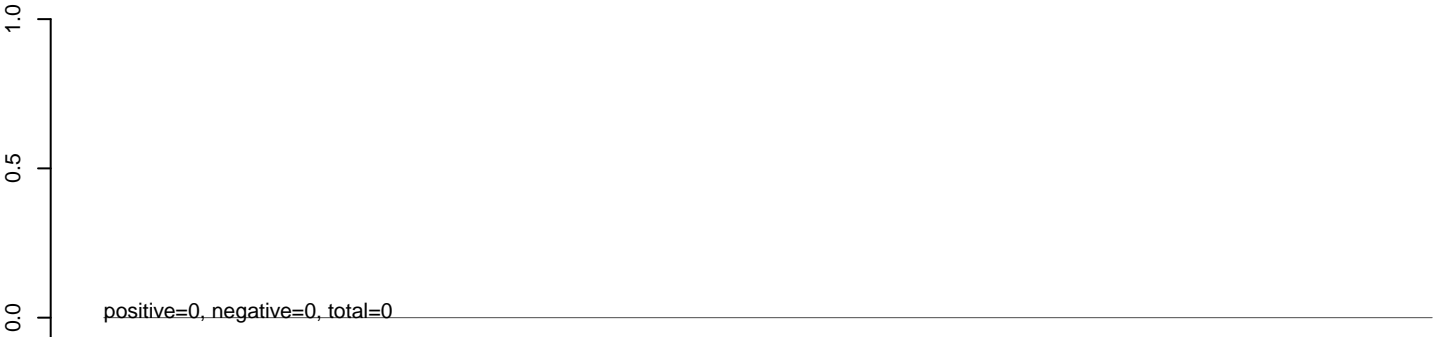


0 500 1000 1500 2000 2500 3000 3500

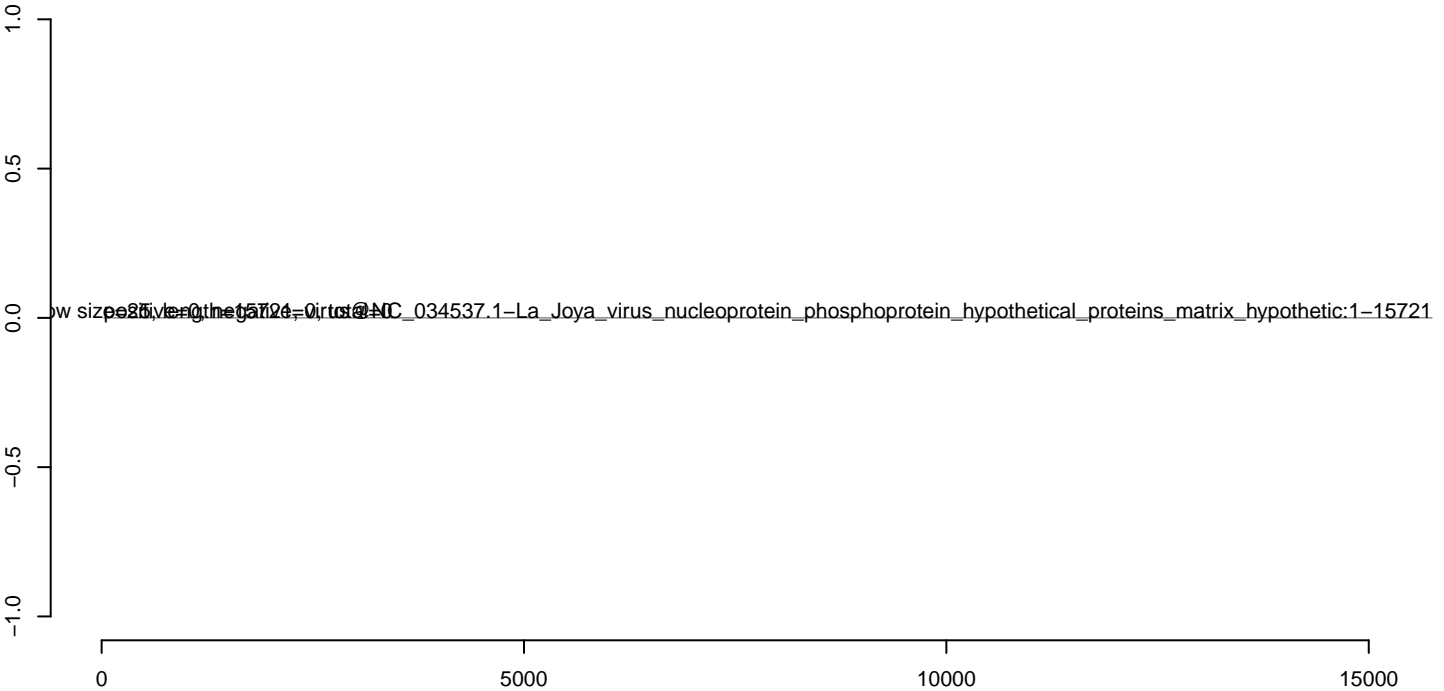
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



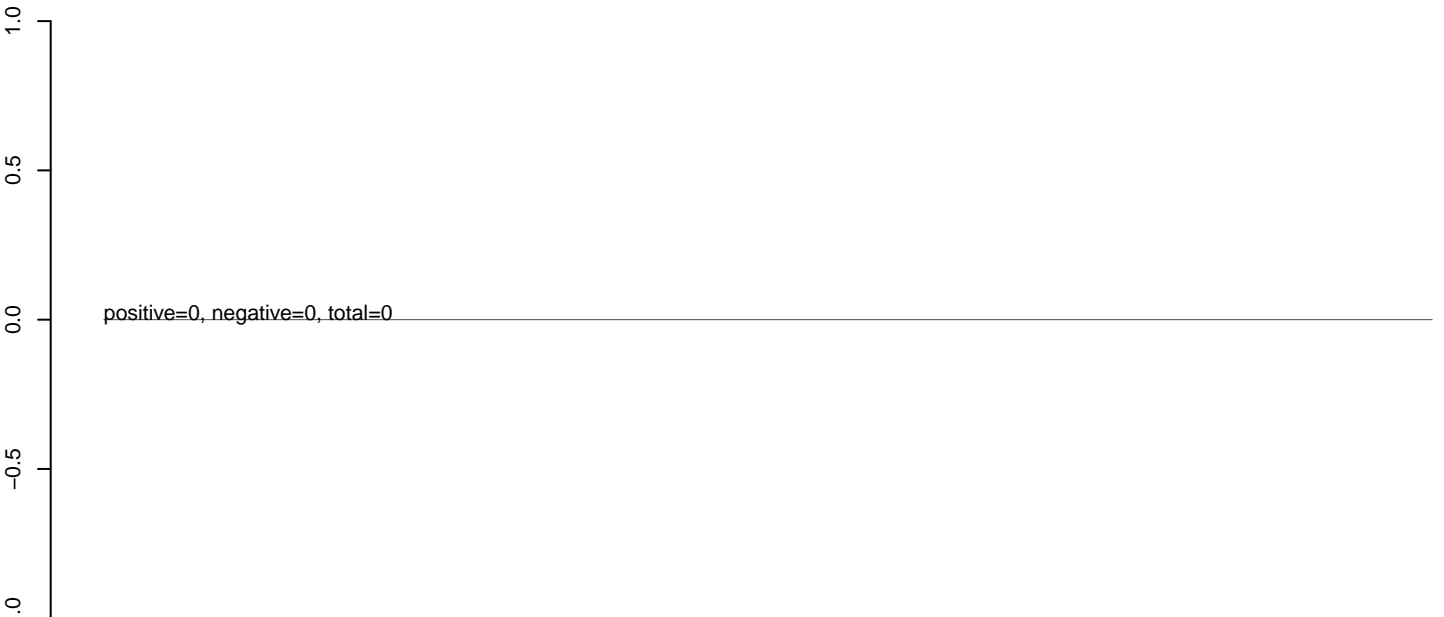
AeAlbo_C636_cells.rep



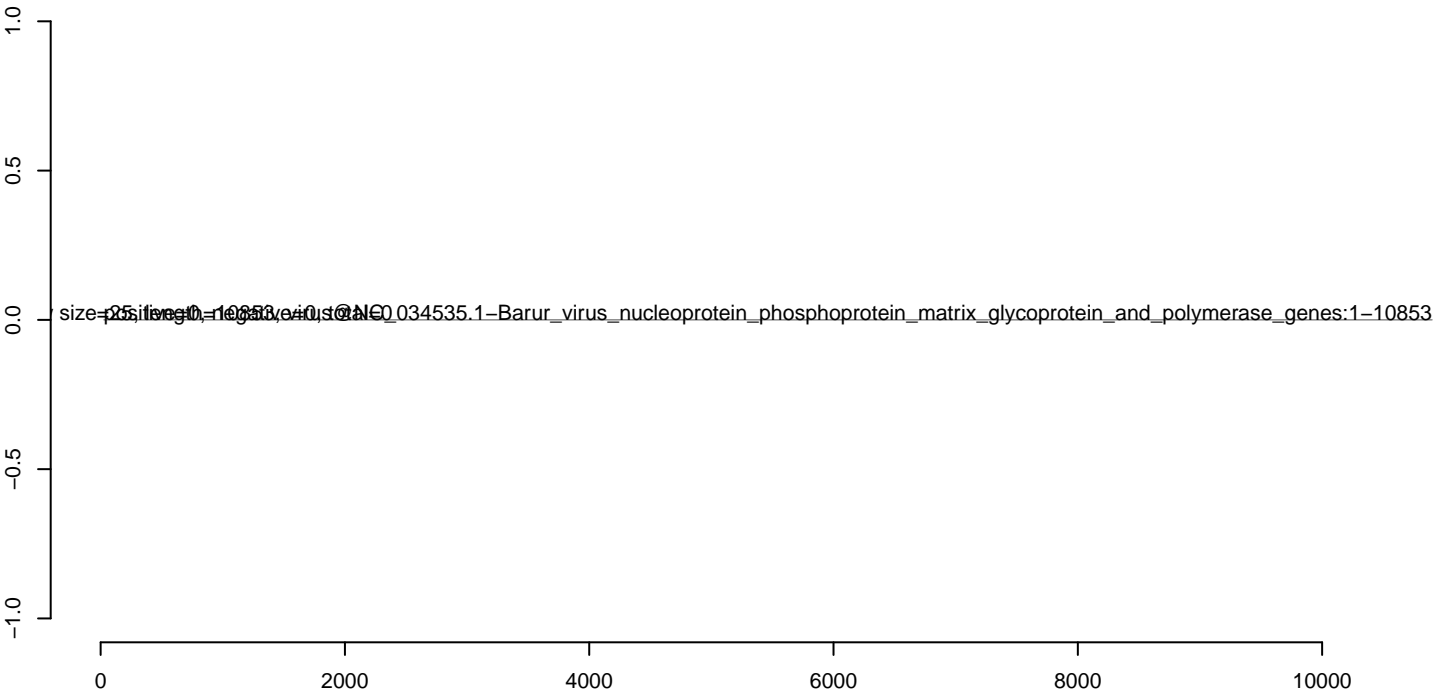
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



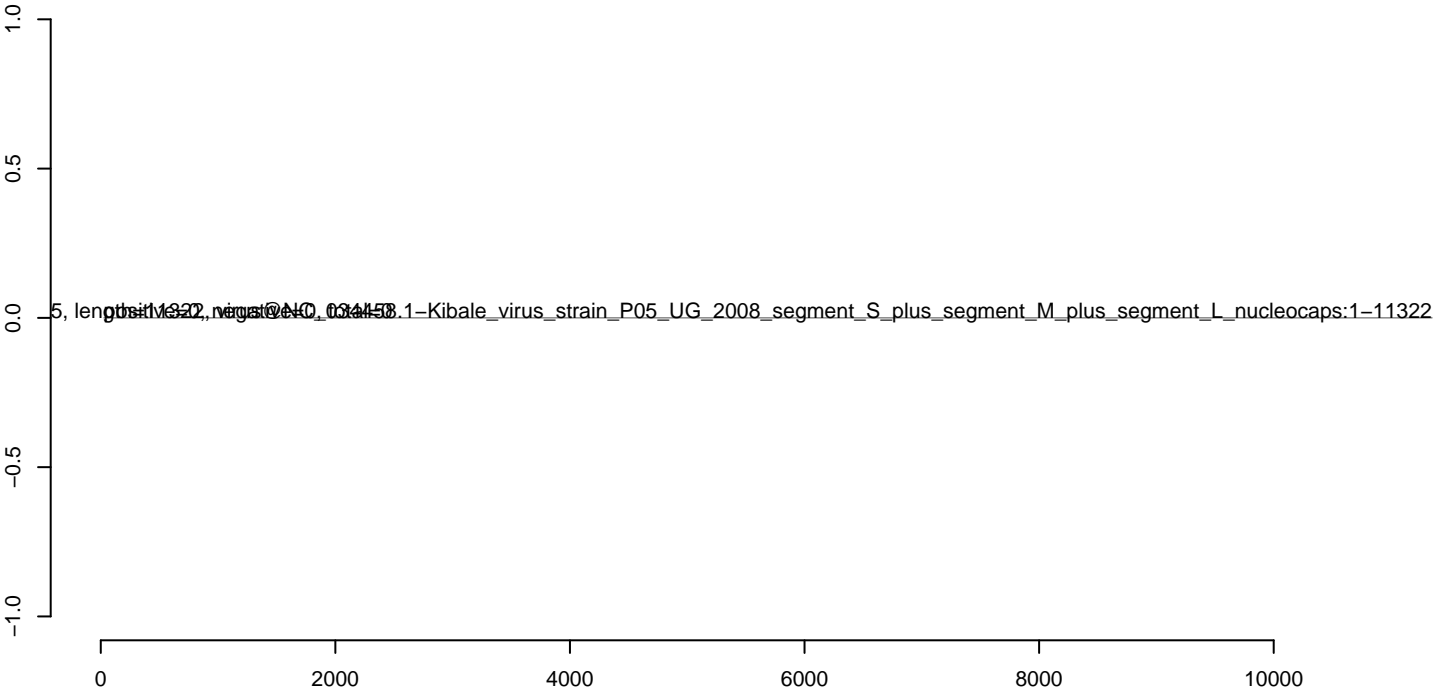
AeAlbo_C636_cells.18_23.rep



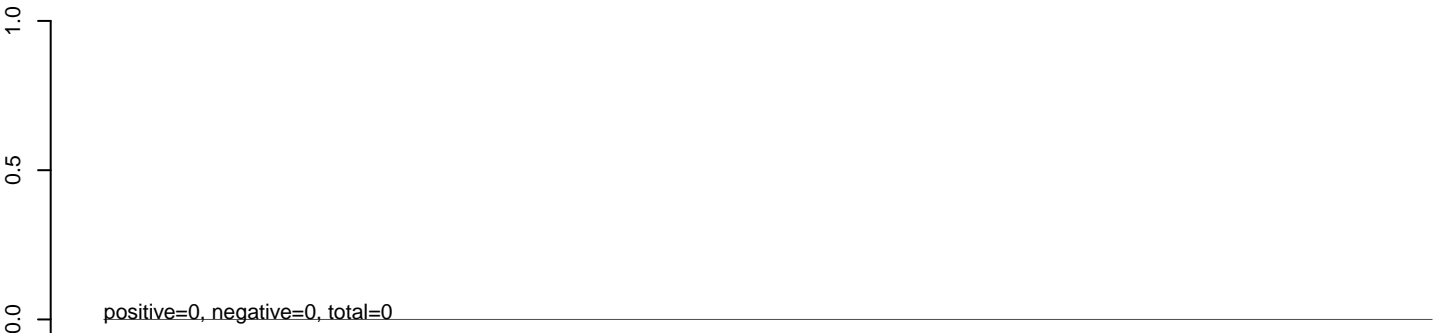
AeAlbo_C636_cells.24_35.rep



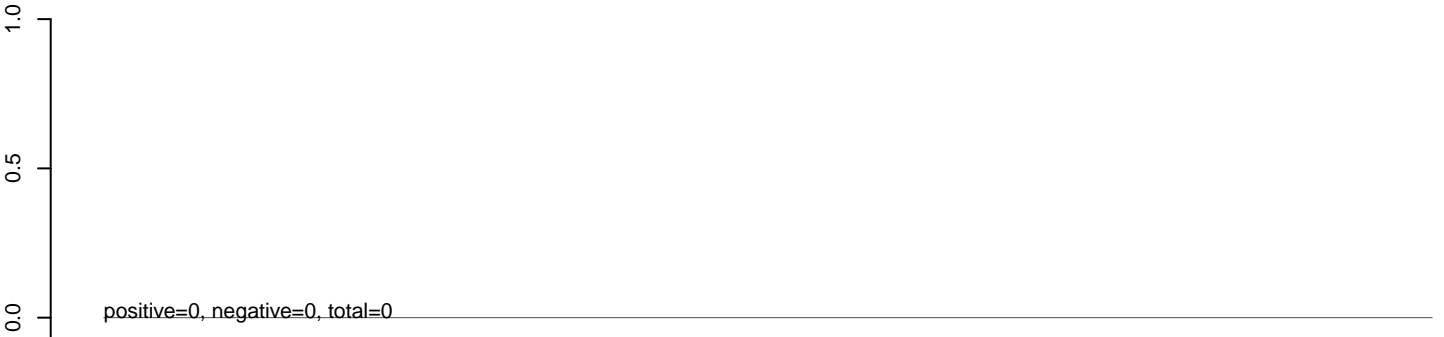
AeAlbo_C636_cells.rep



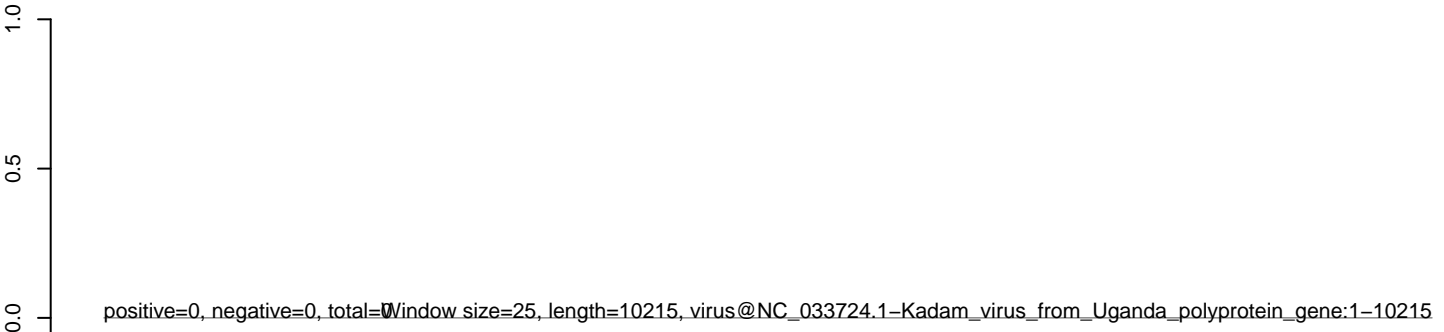
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



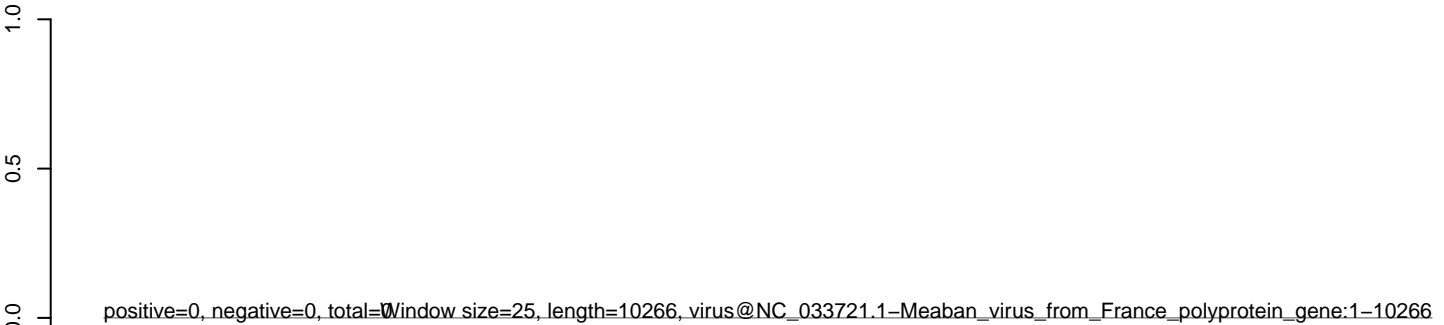
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



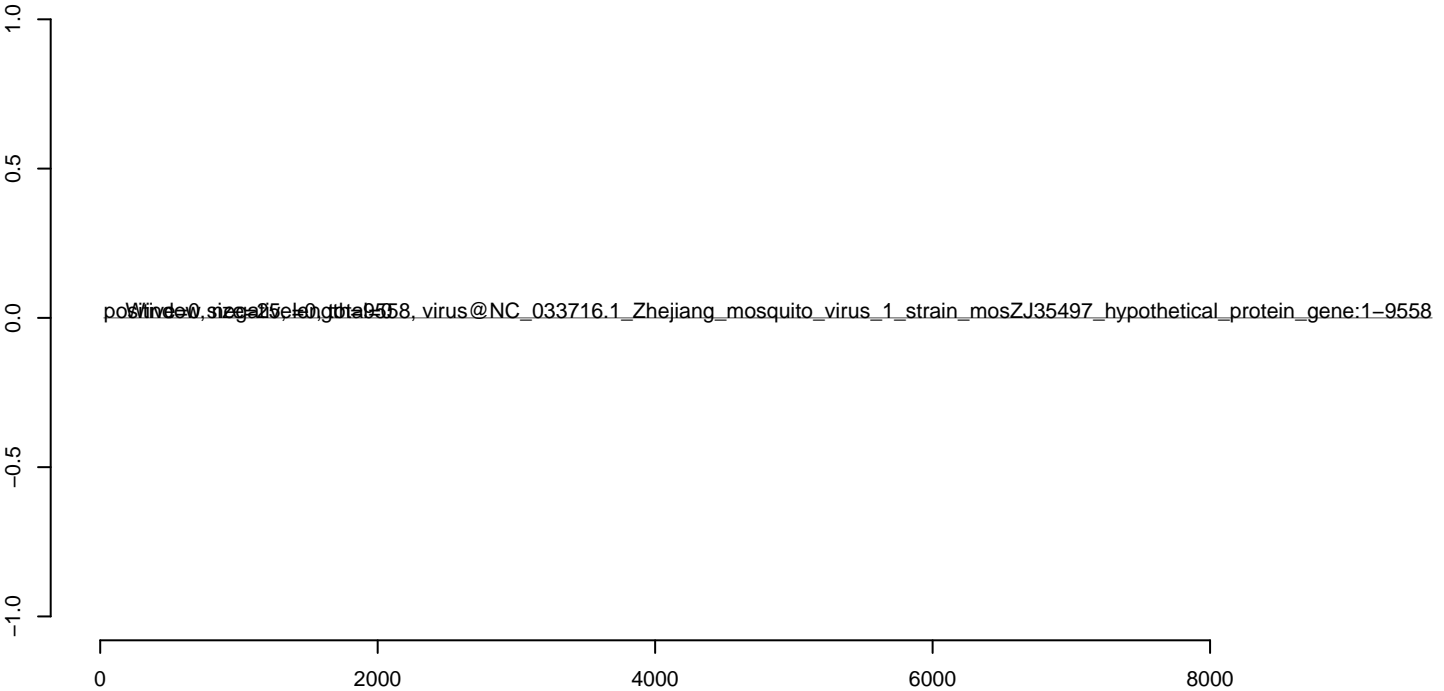
AeAlbo_C636_cells.18_23.rep



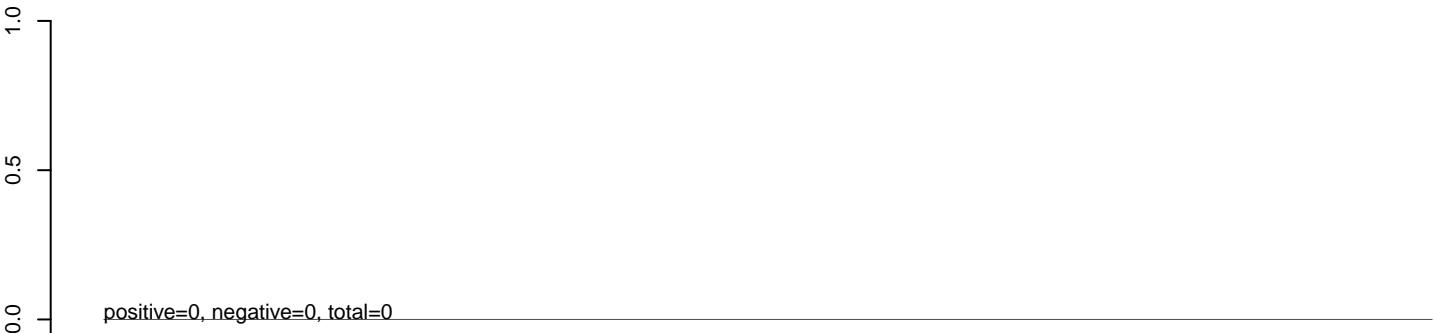
AeAlbo_C636_cells.24_35.rep



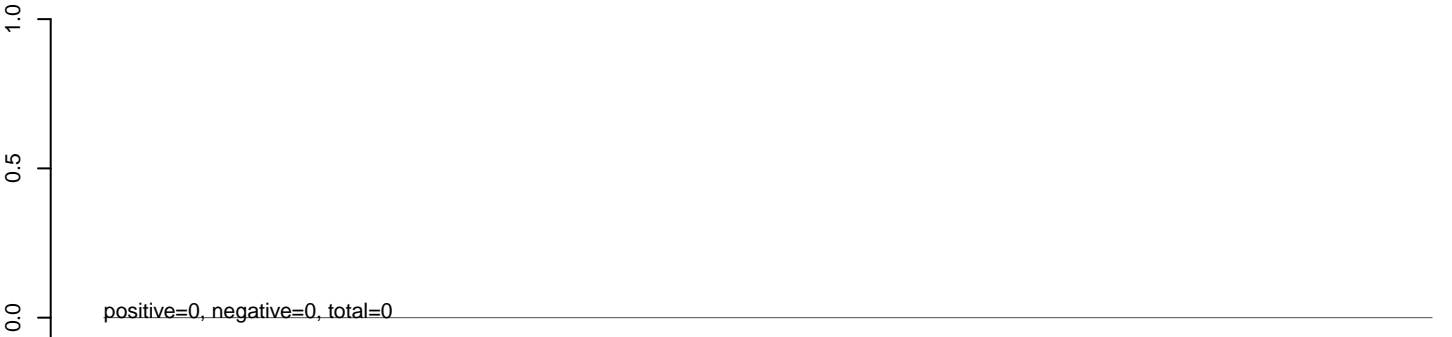
AeAlbo_C636_cells.rep



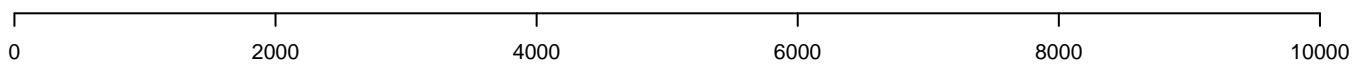
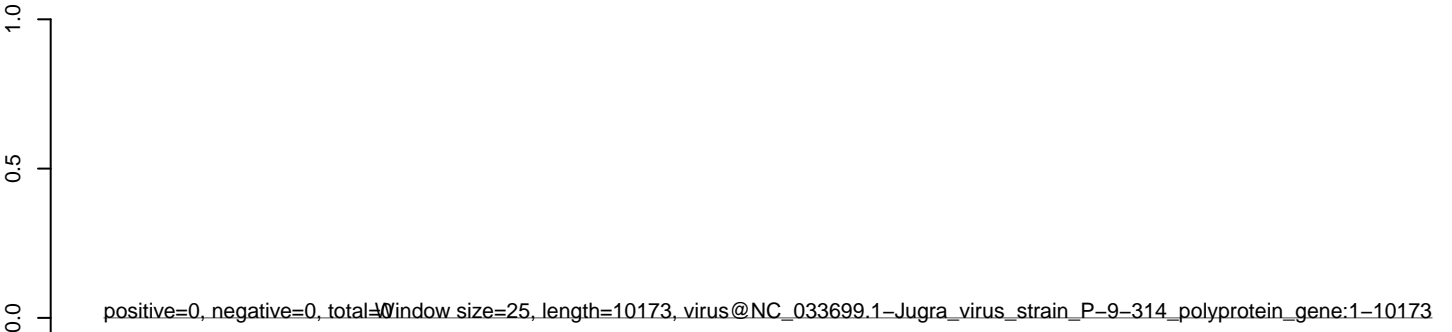
AeAlbo_C636_cells.18_23.rep



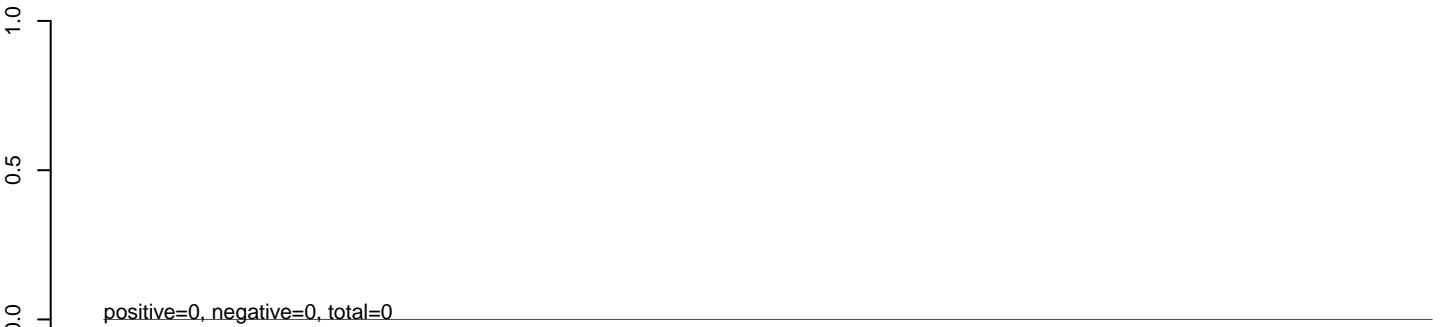
AeAlbo_C636_cells.24_35.rep



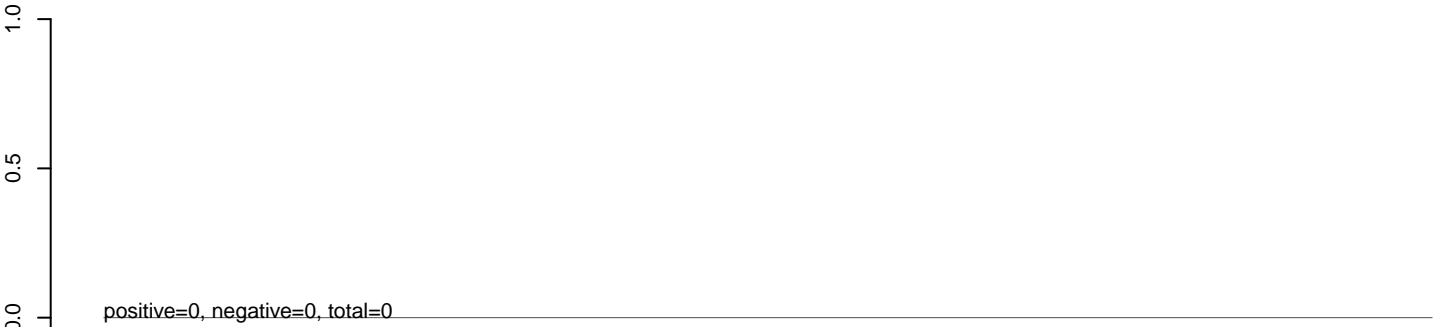
AeAlbo_C636_cells.rep



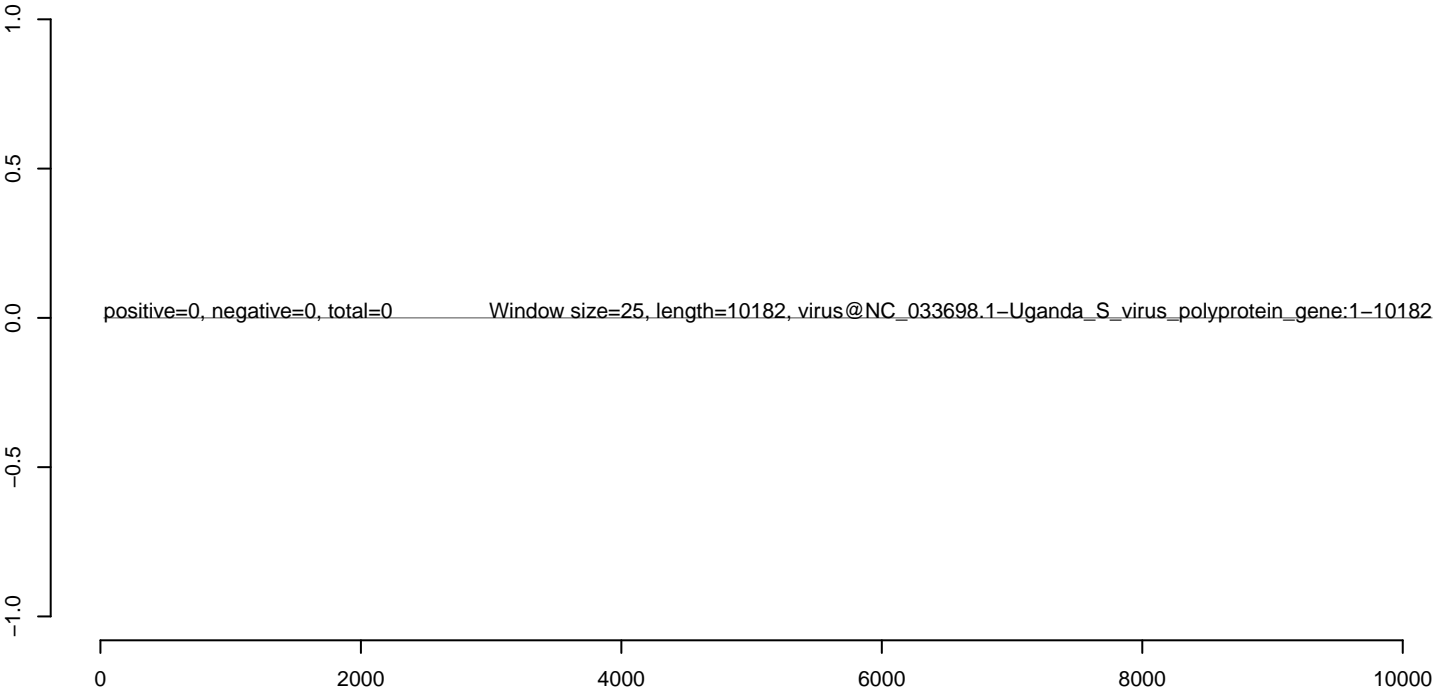
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



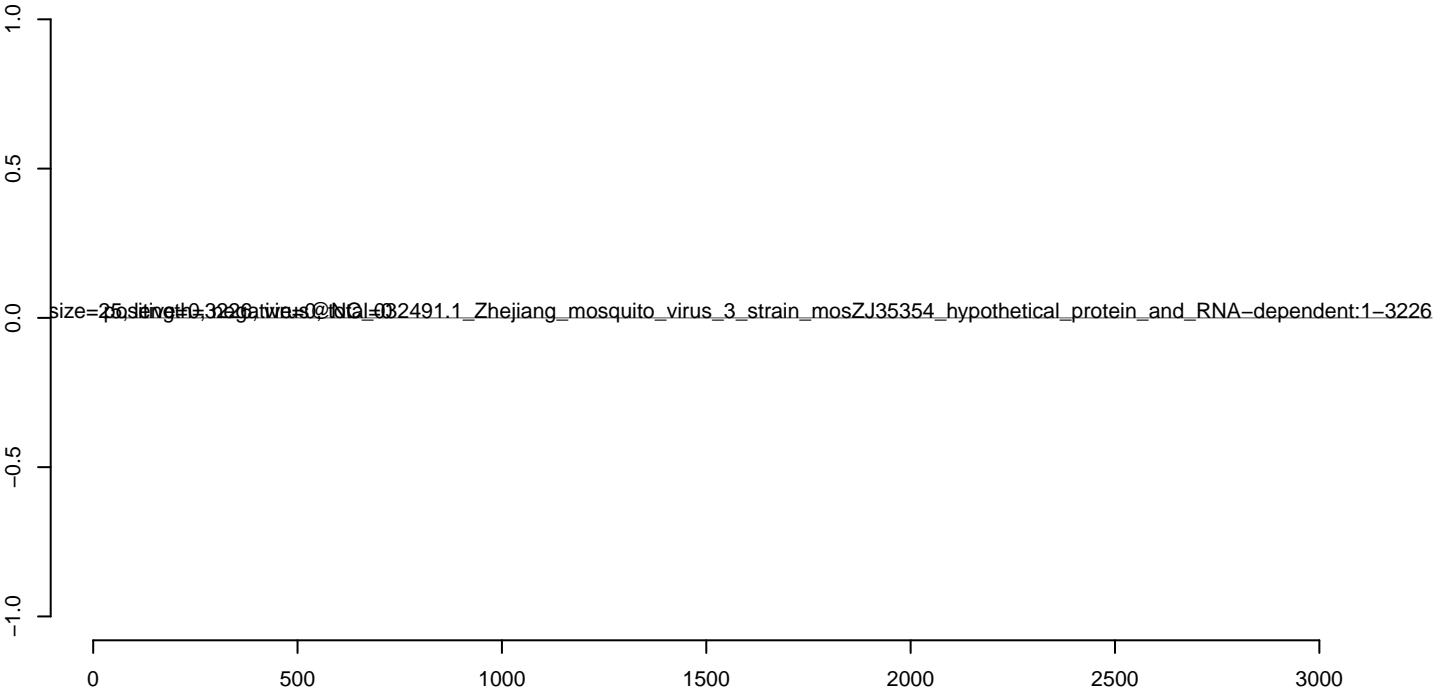
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



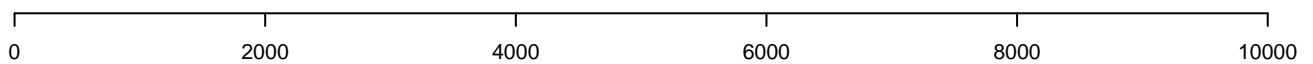
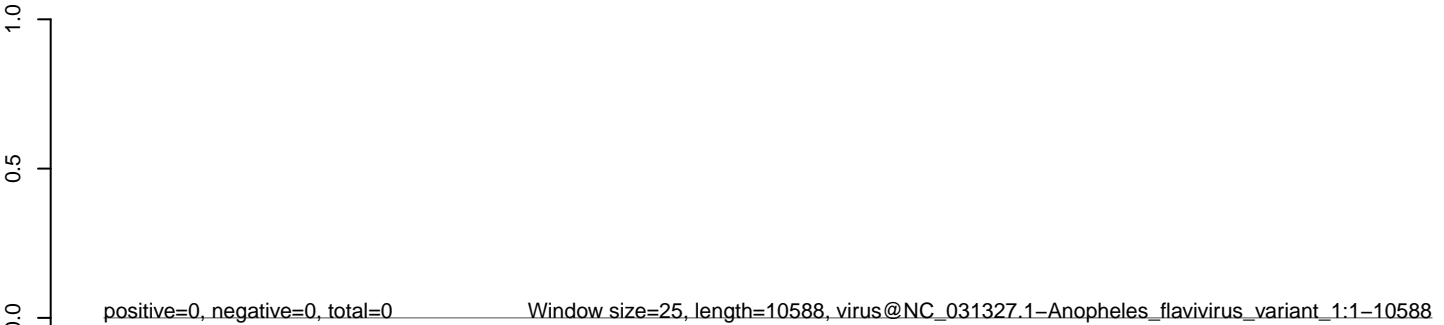
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



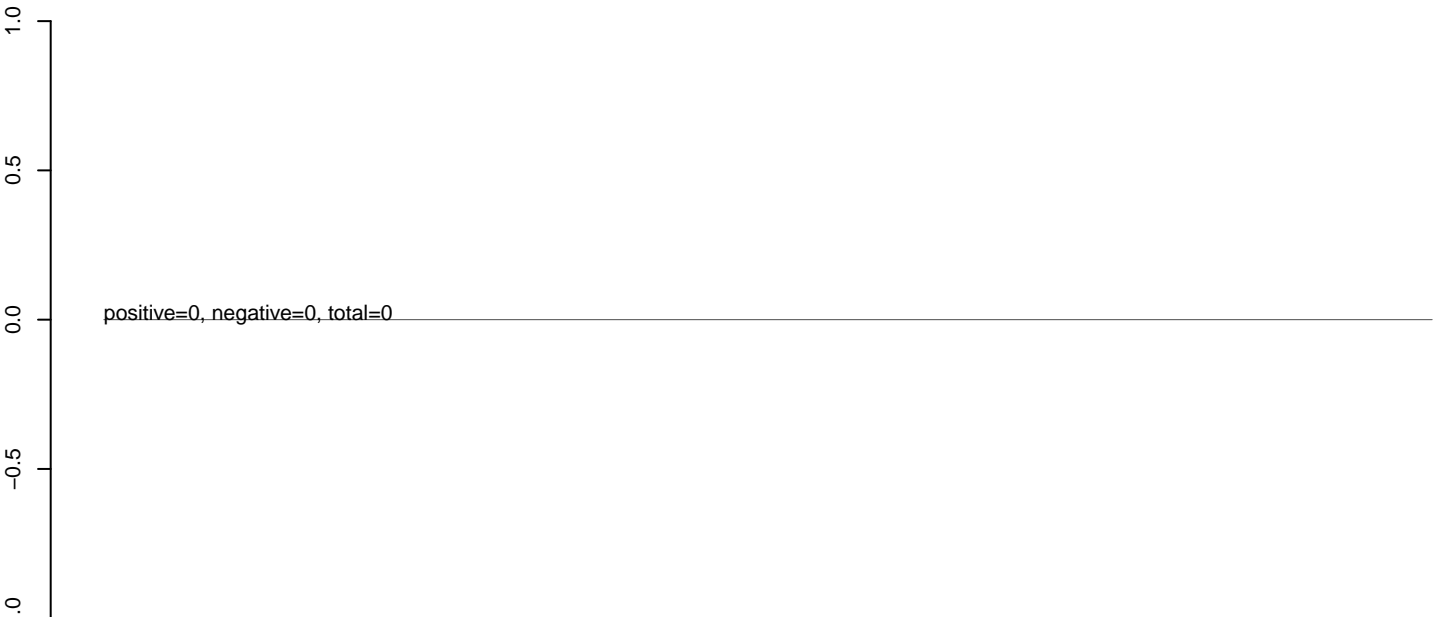
AeAlbo_C636_cells.rep



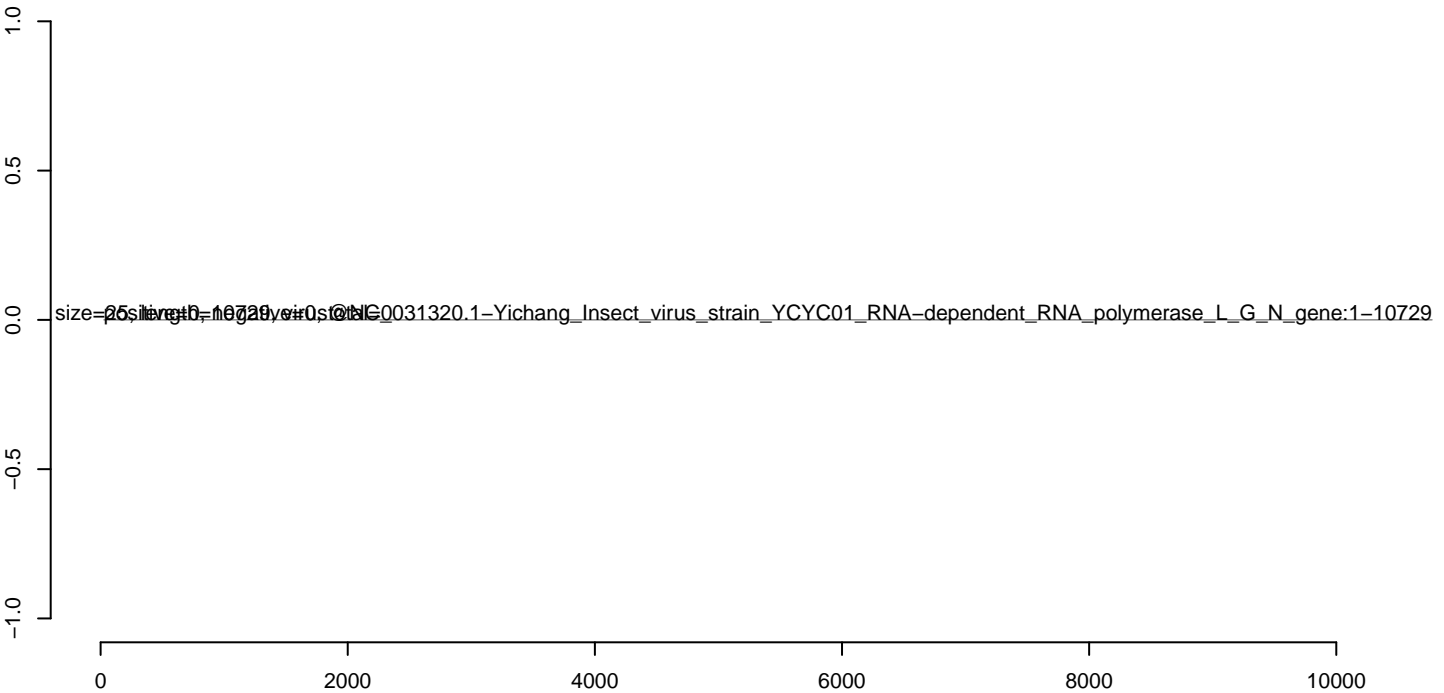
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



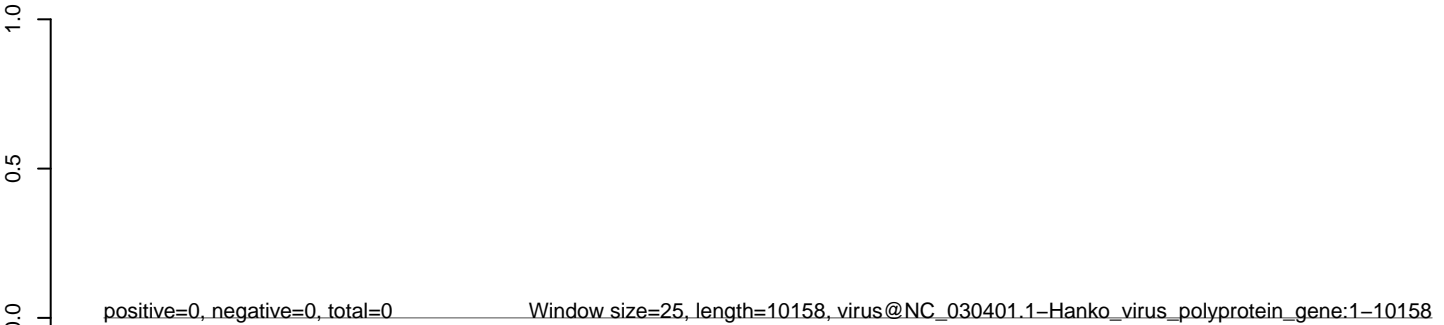
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

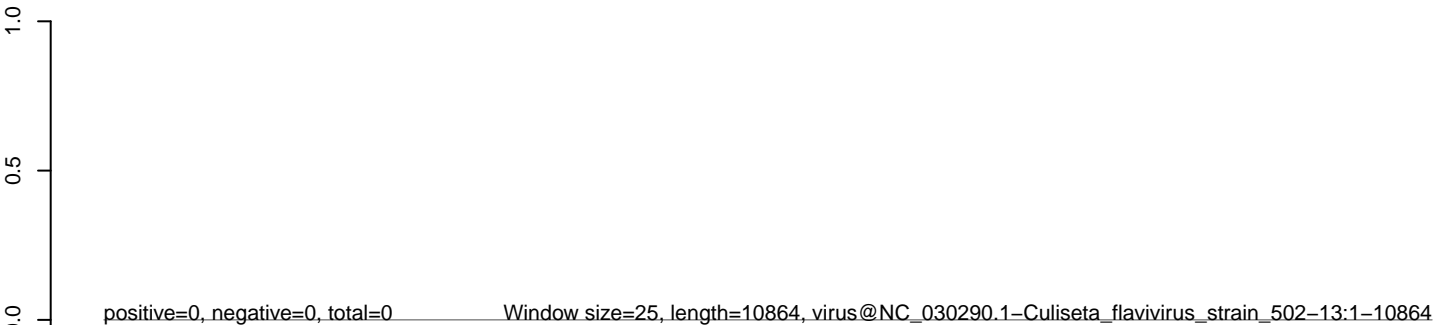
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

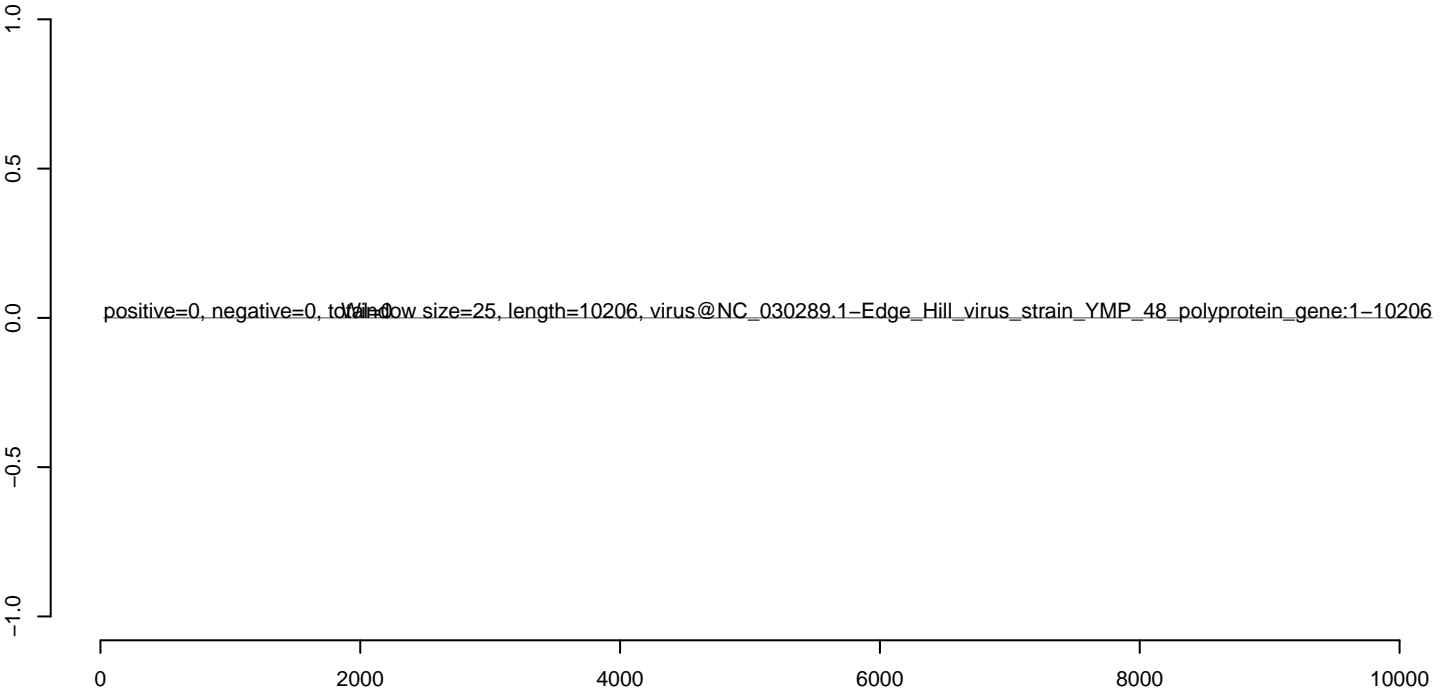
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



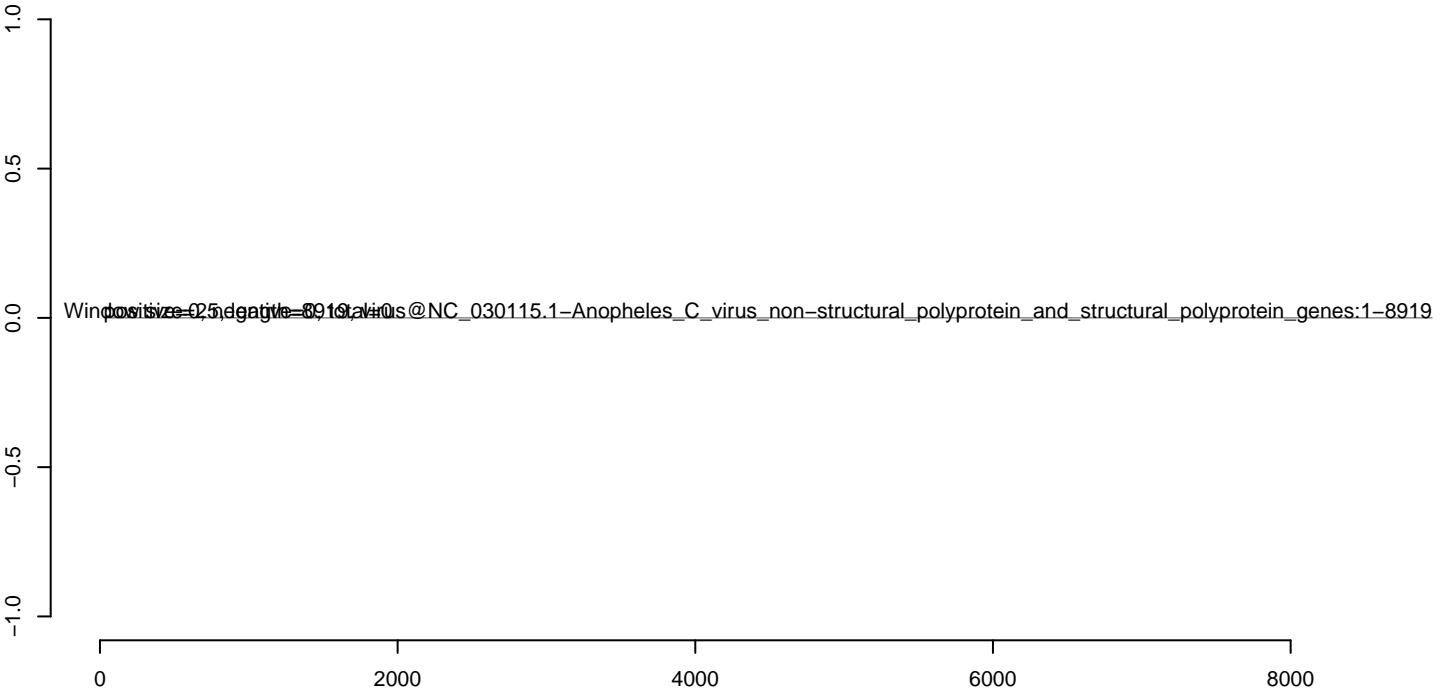
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



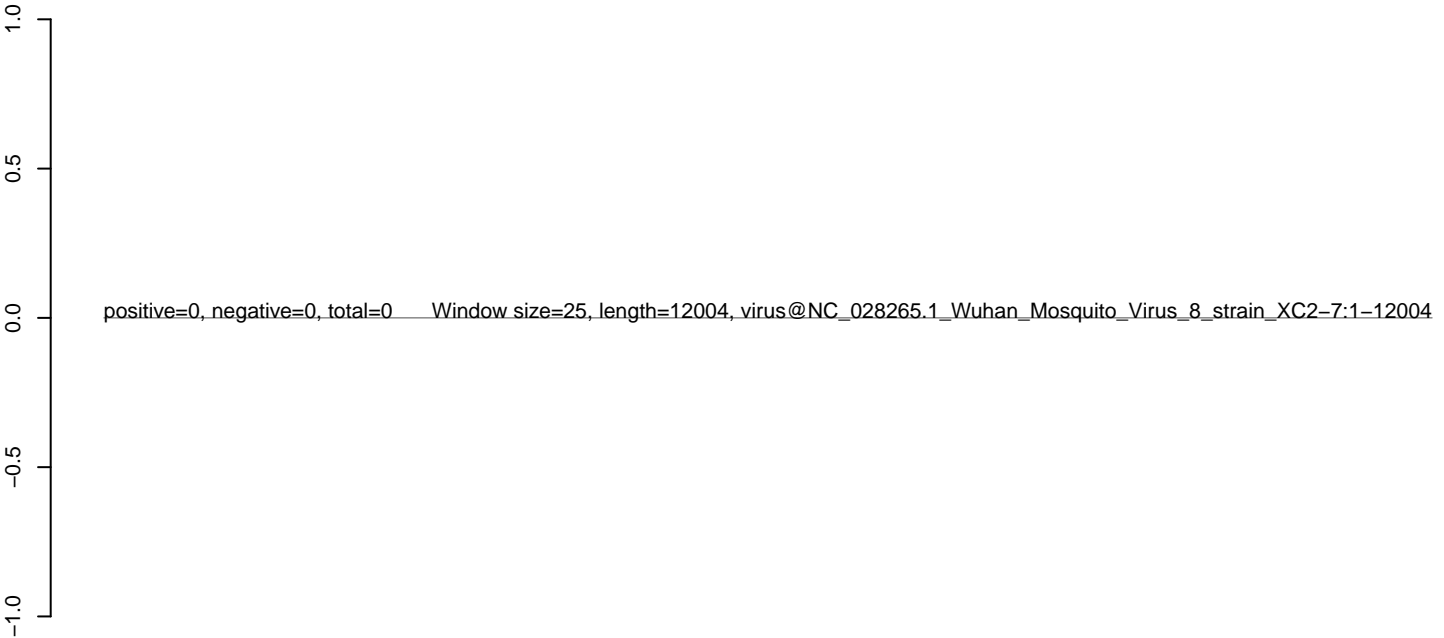
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



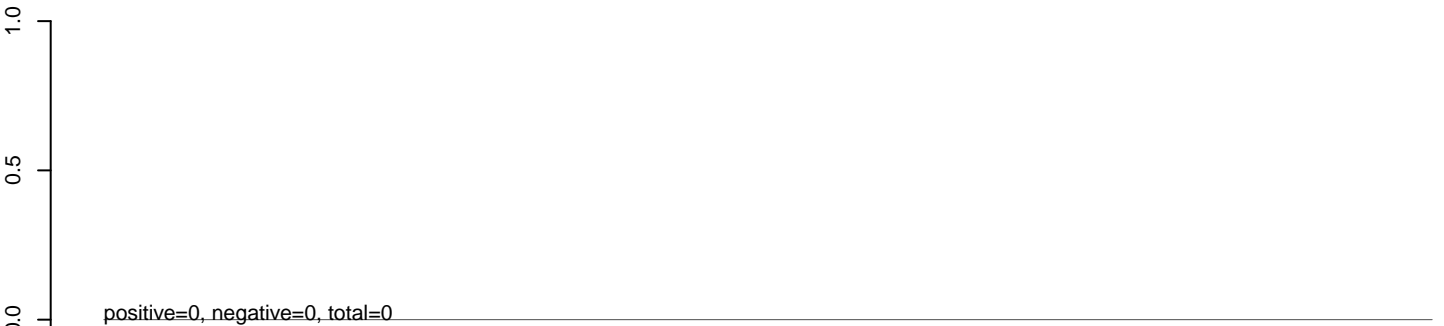
AeAlbo_C636_cells.rep



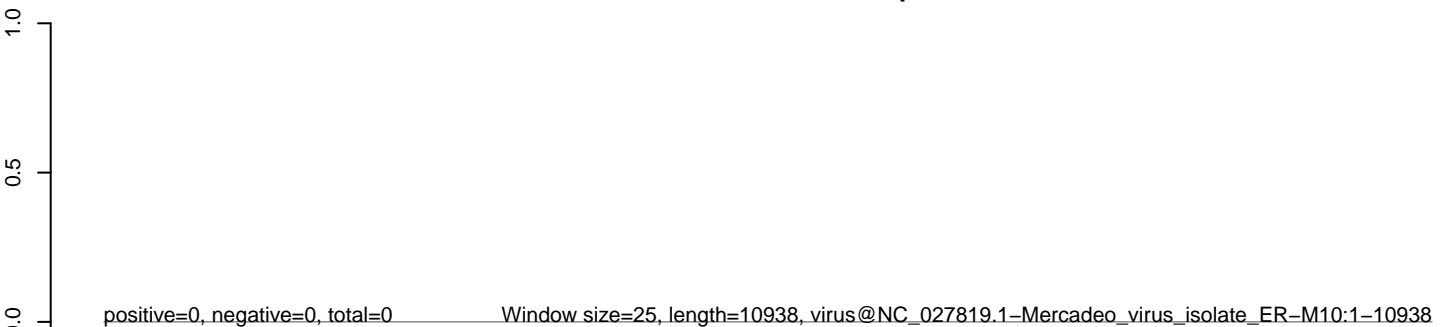
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



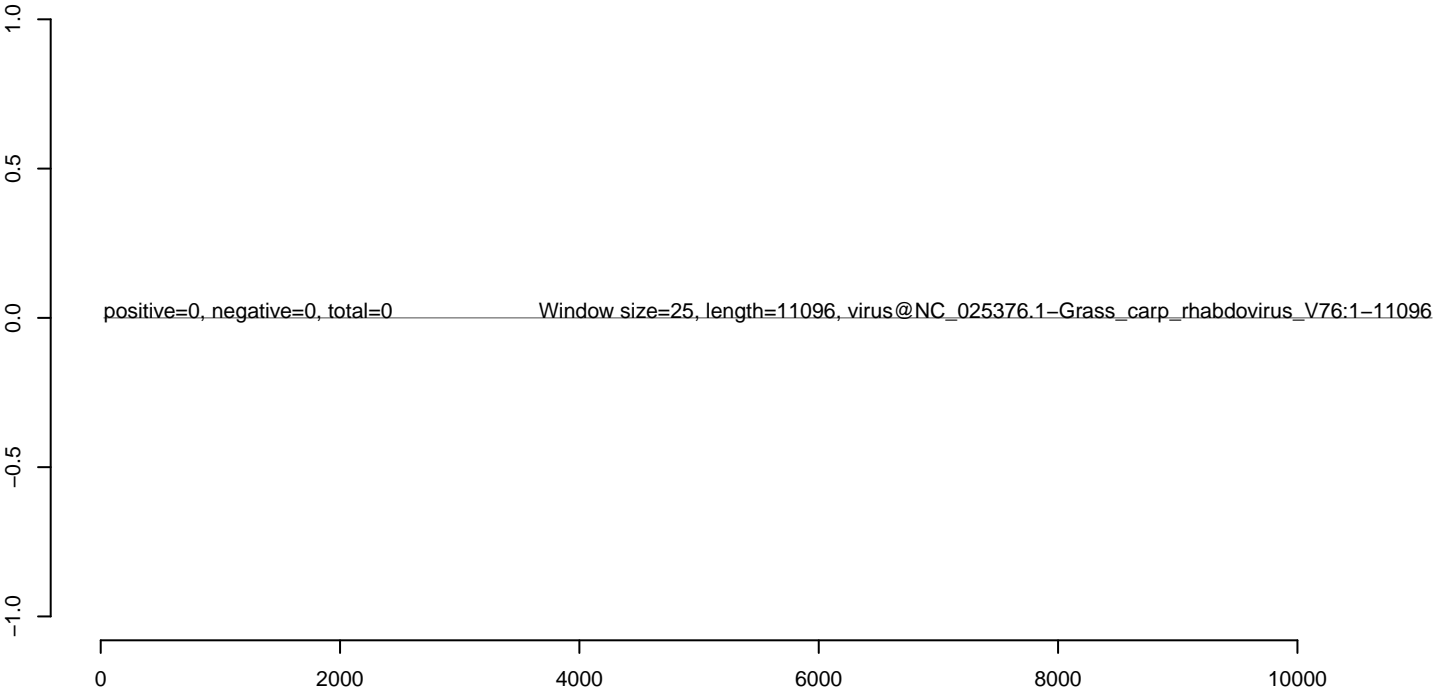
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



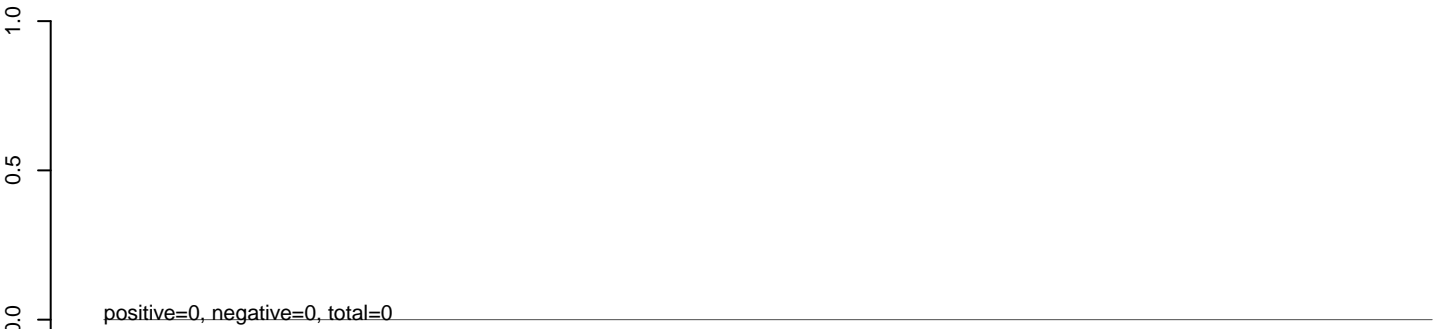
AeAlbo_C636_cells.rep



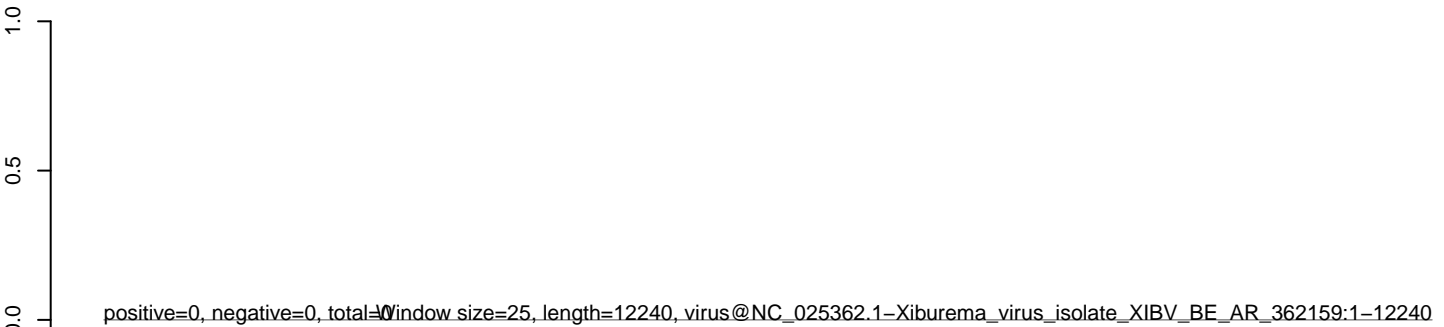
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



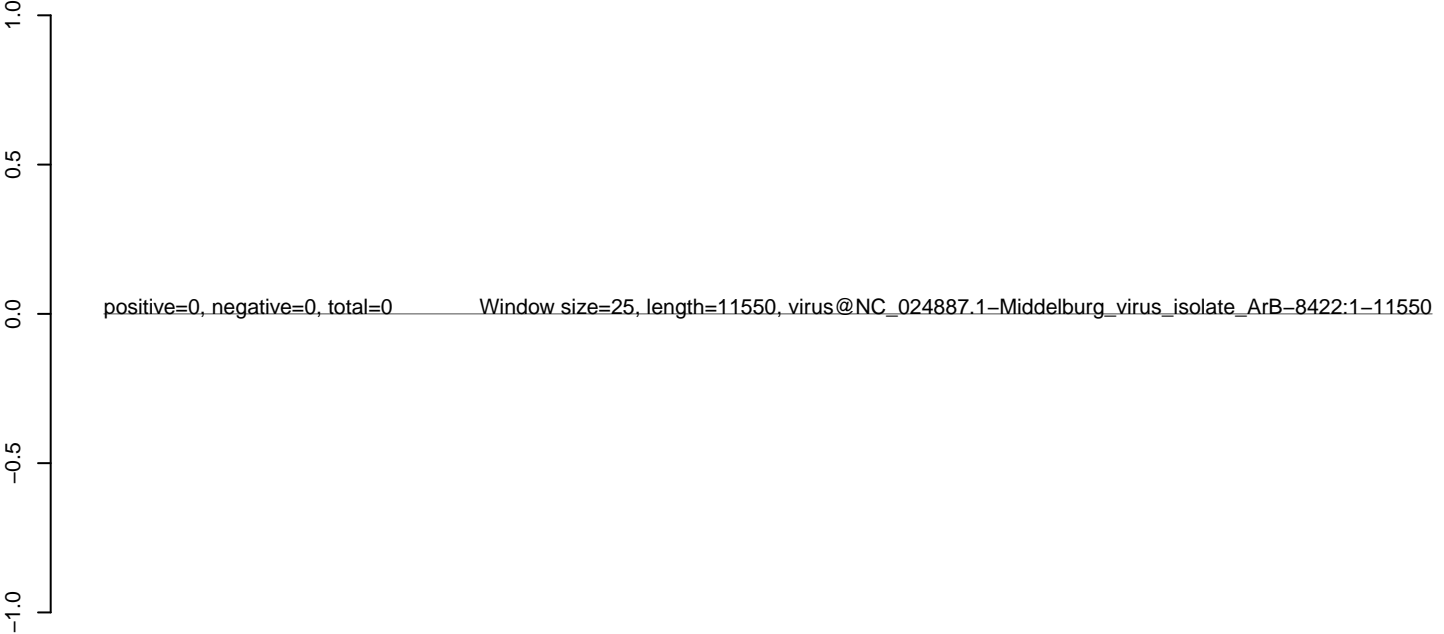
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

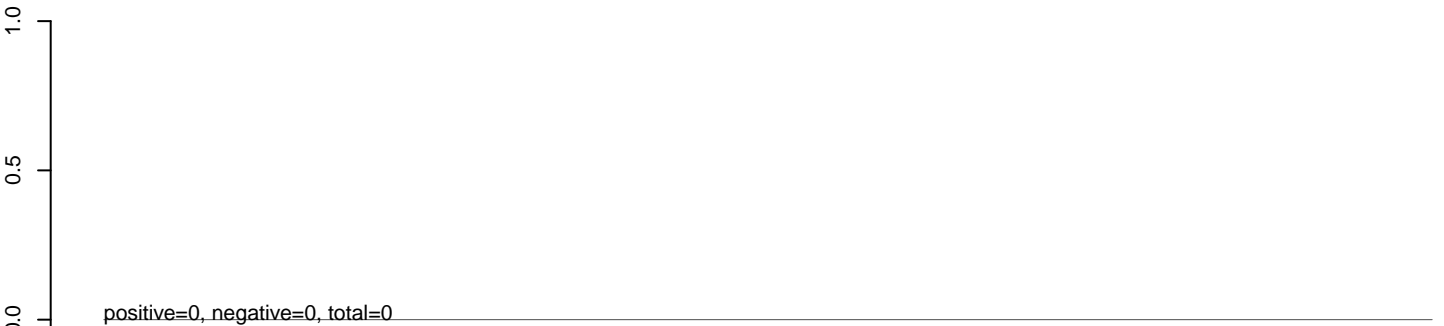


0 2000 4000 6000 8000 10000 12000

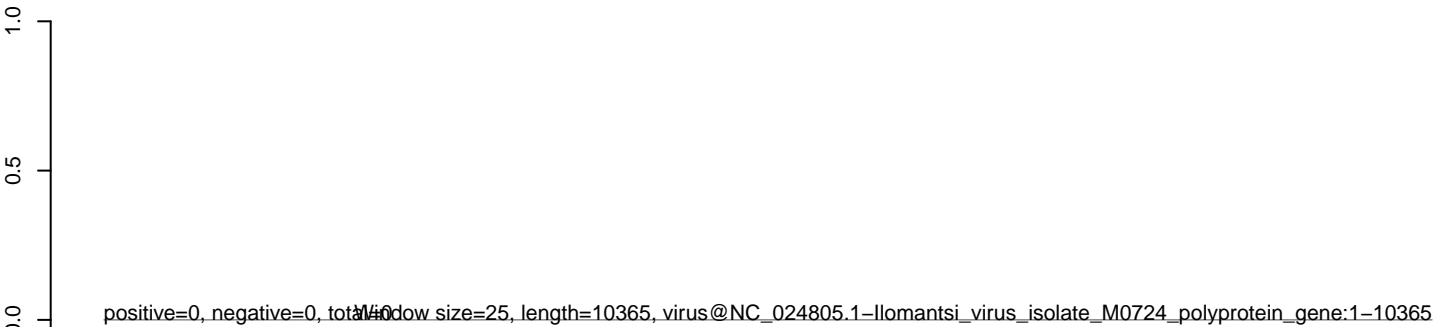
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

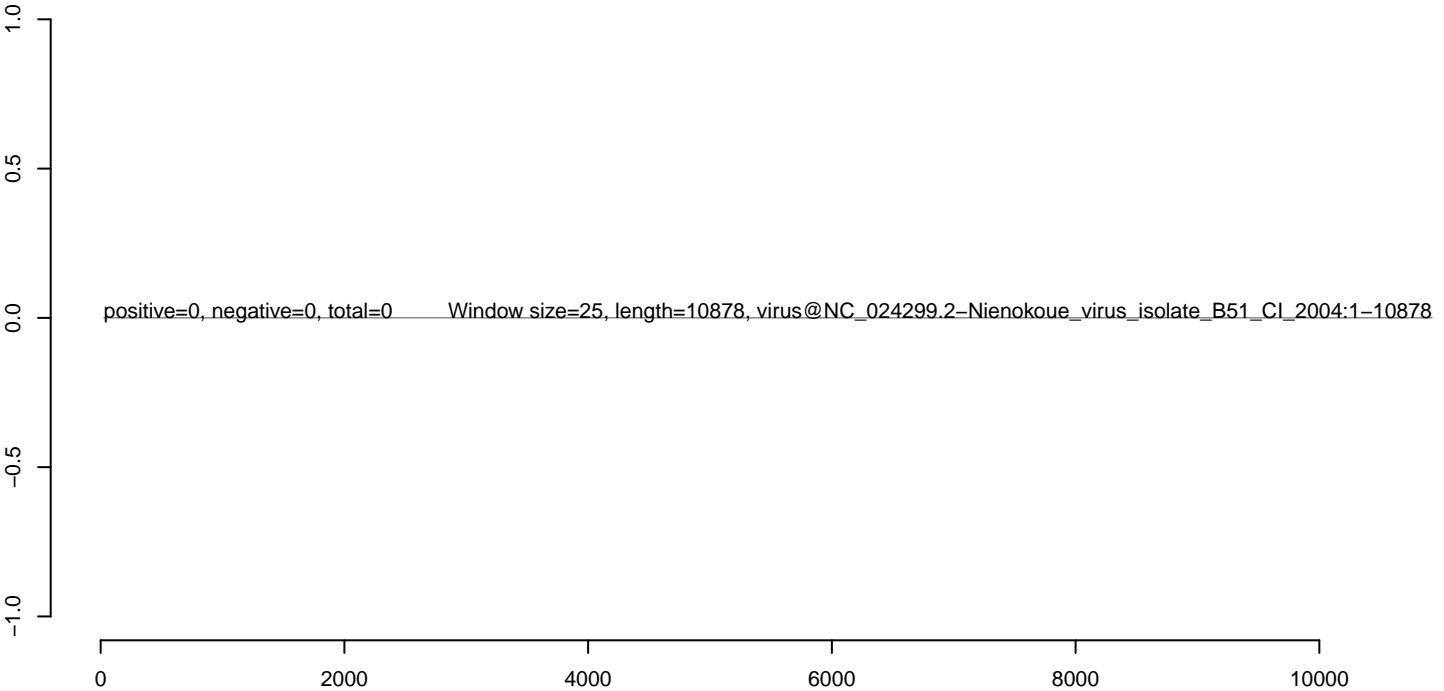
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



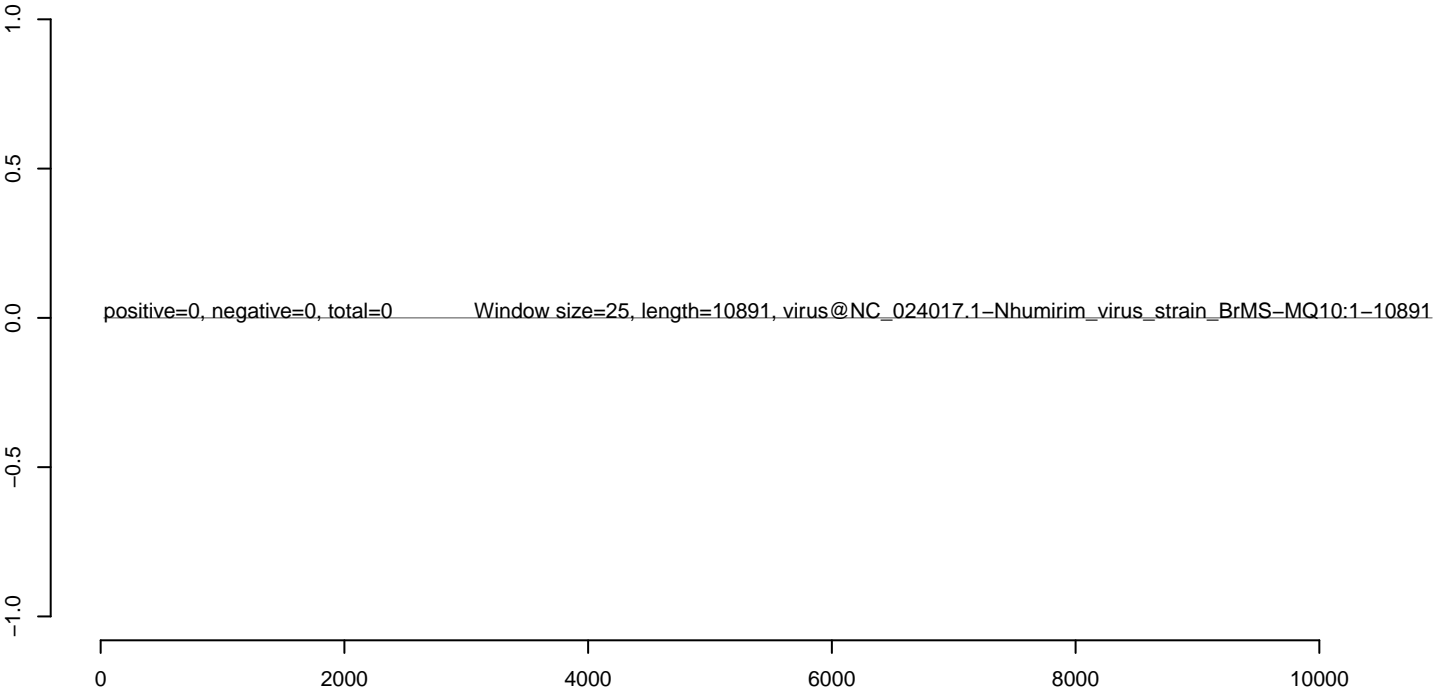
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



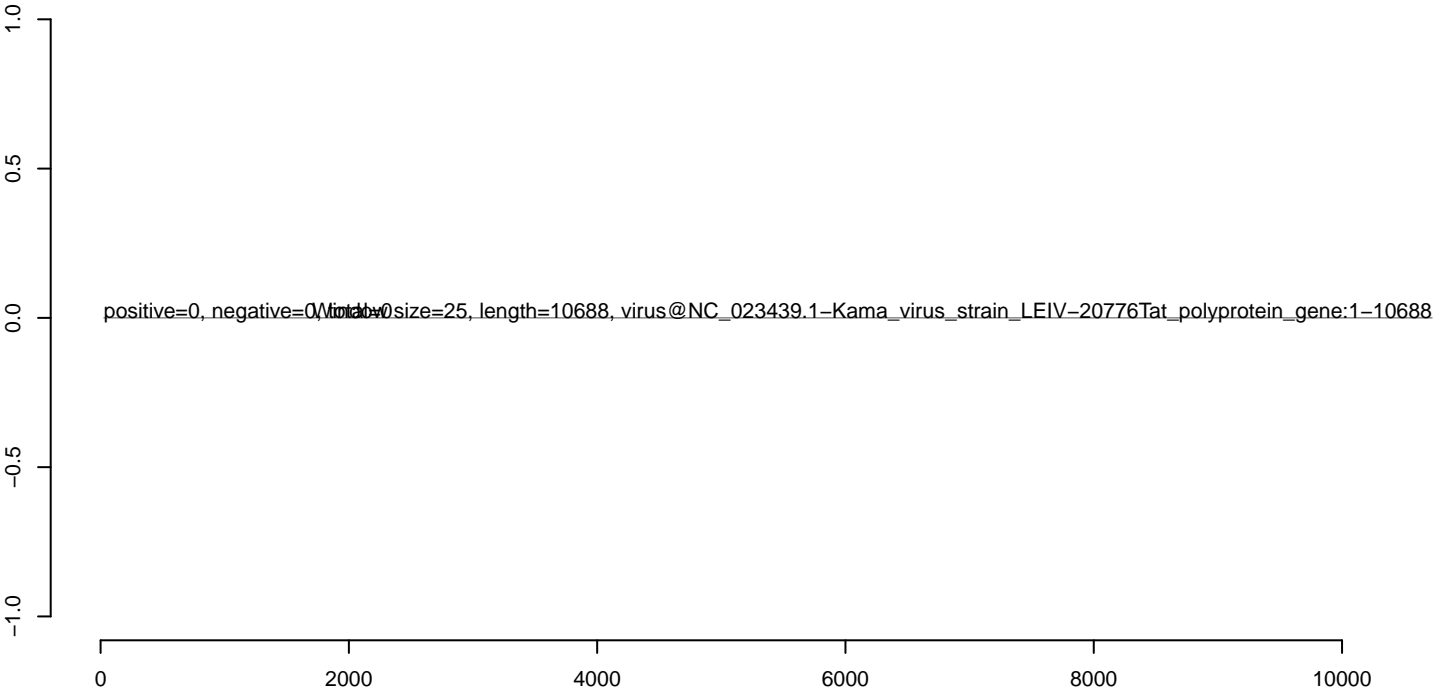
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



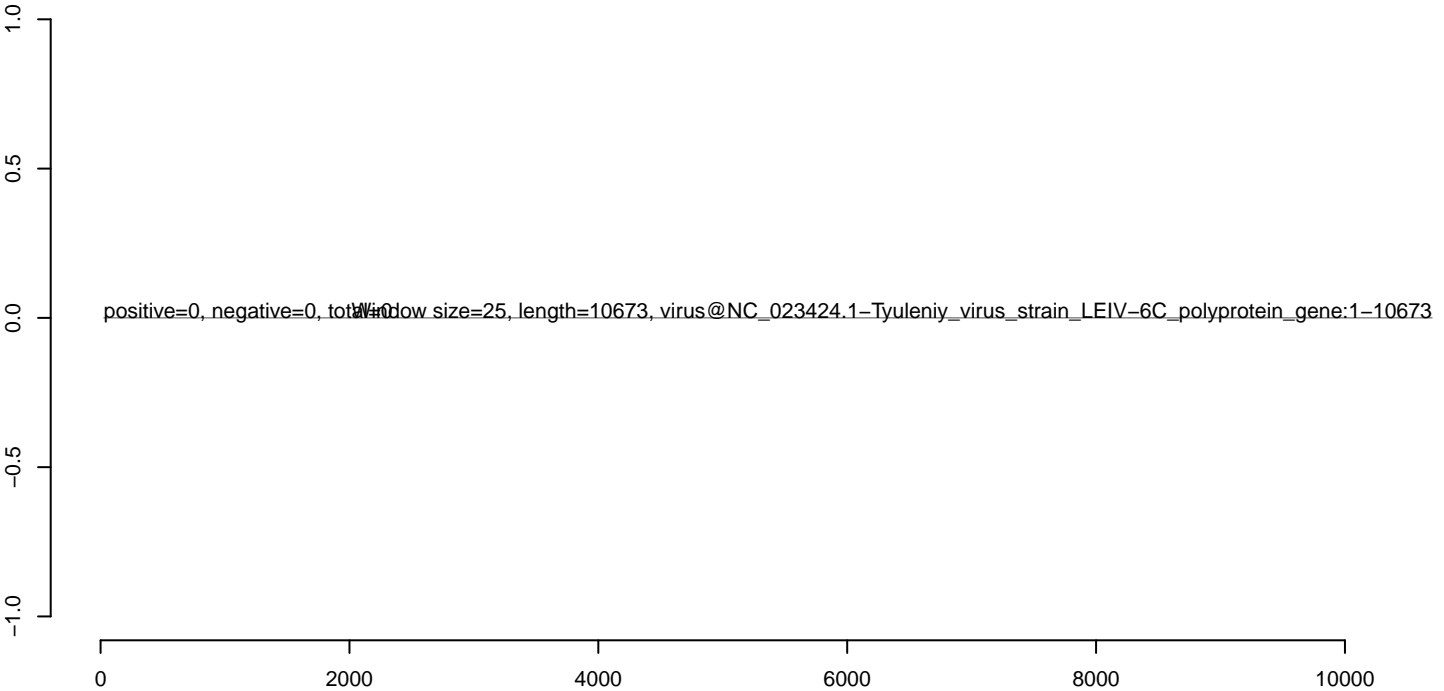
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



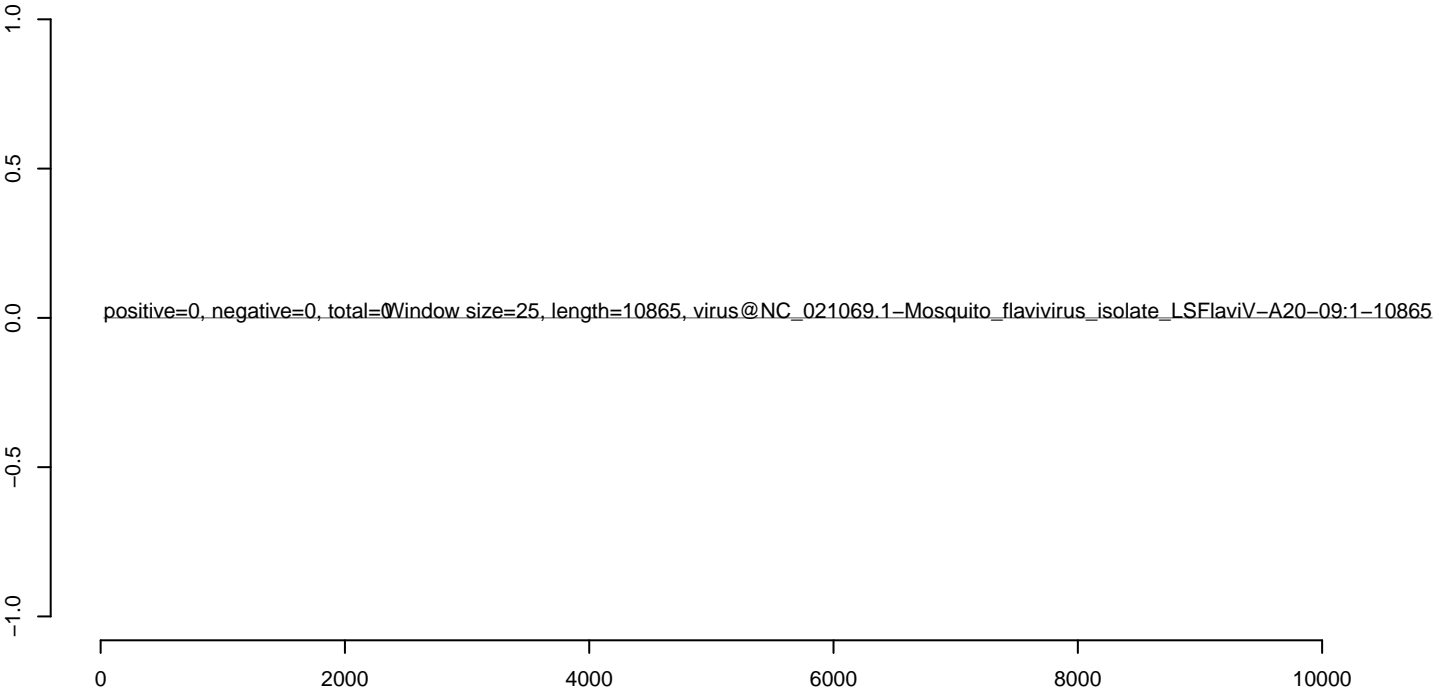
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



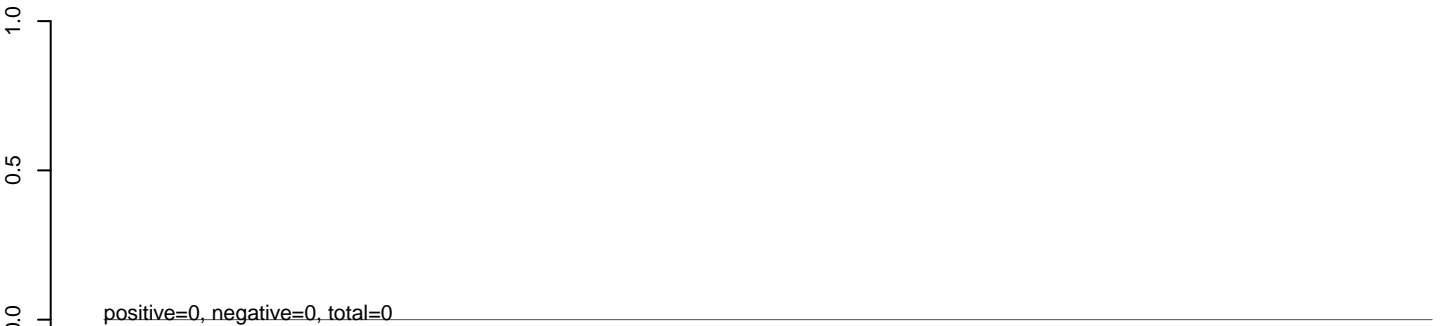
AeAlbo_C636_cells.rep



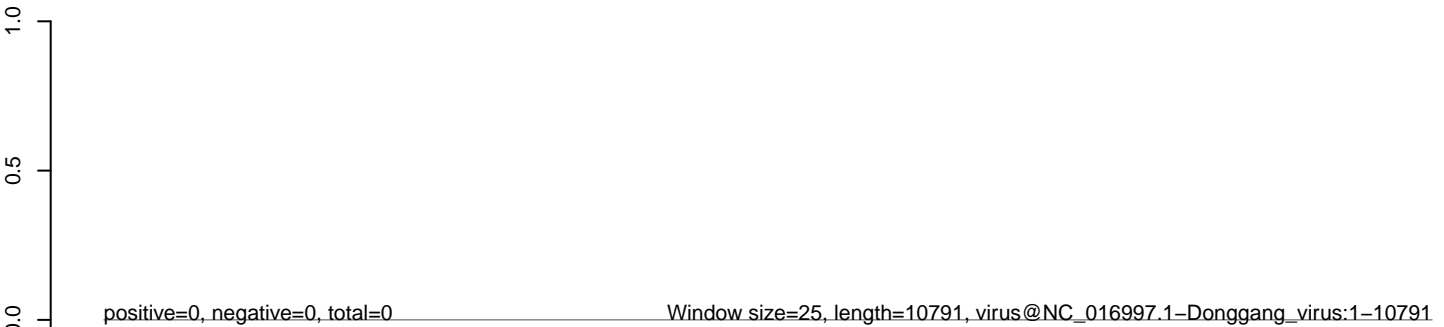
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

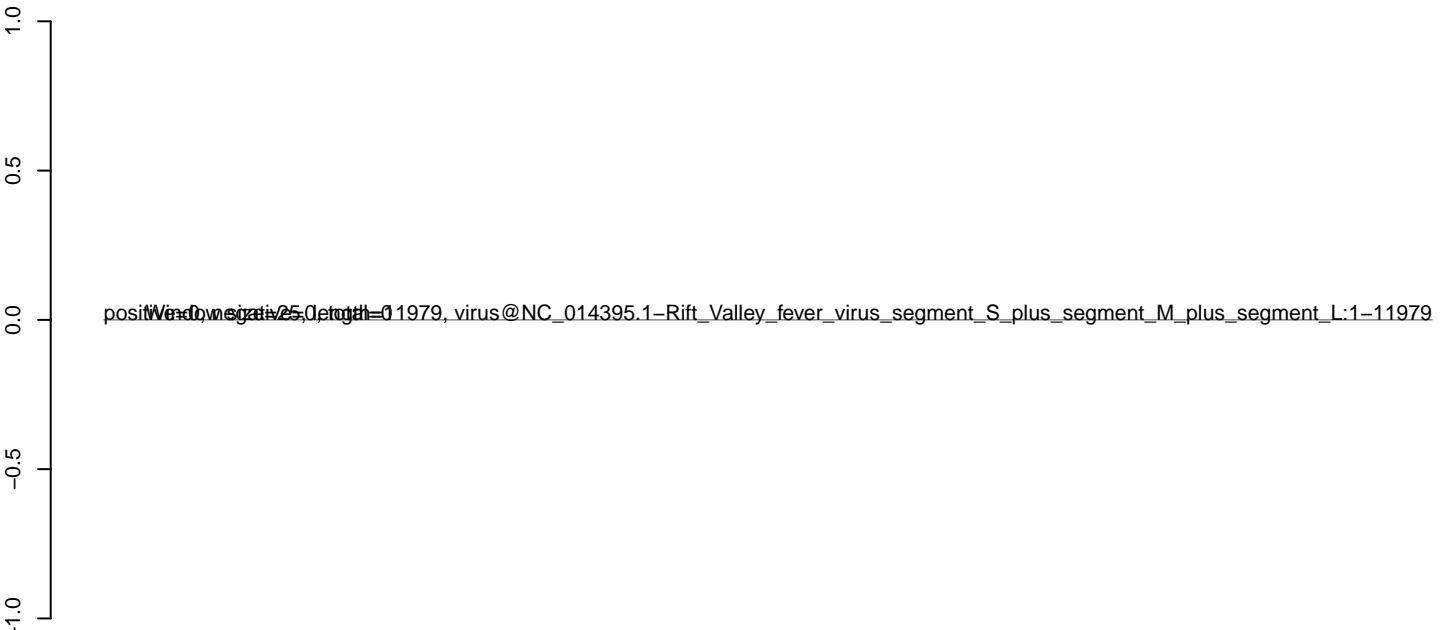
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



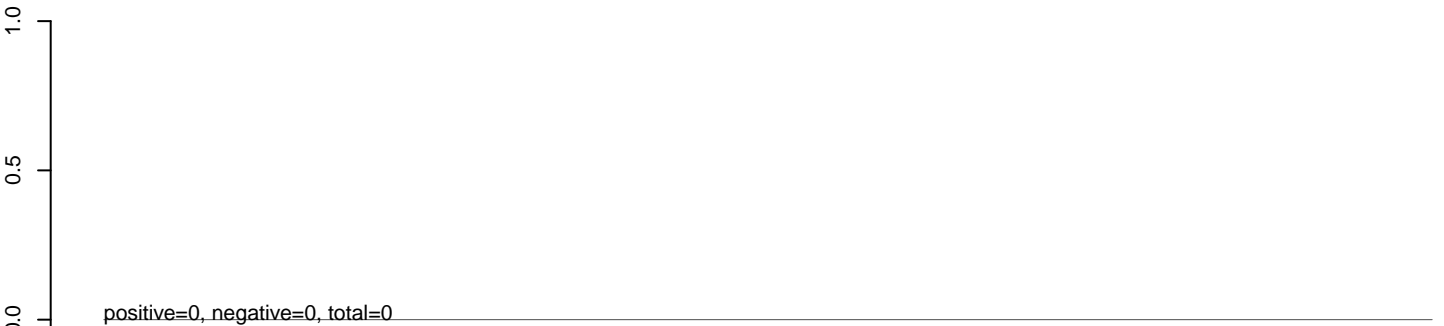
AeAlbo_C636_cells.rep



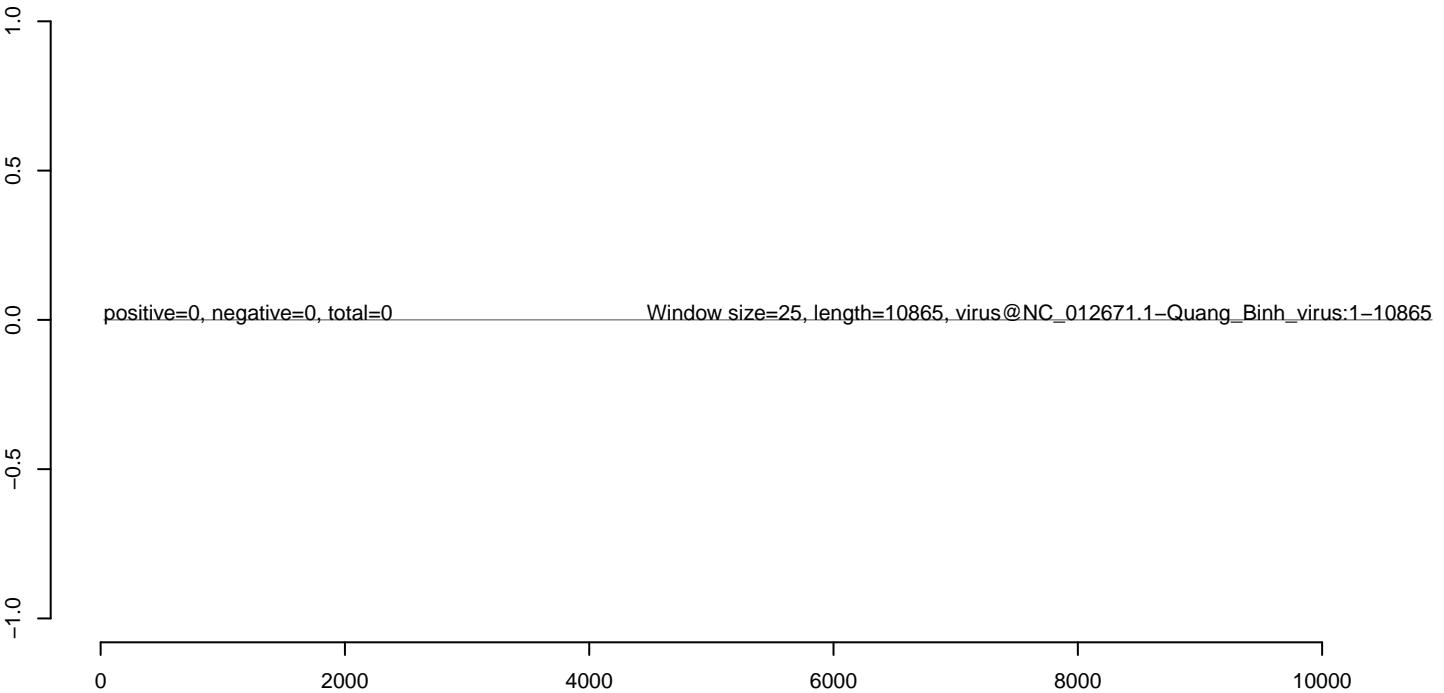
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



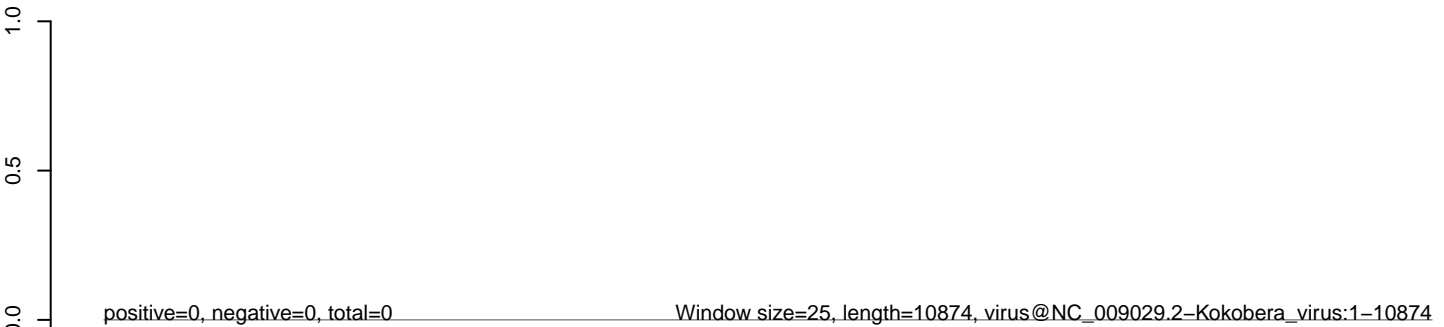
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

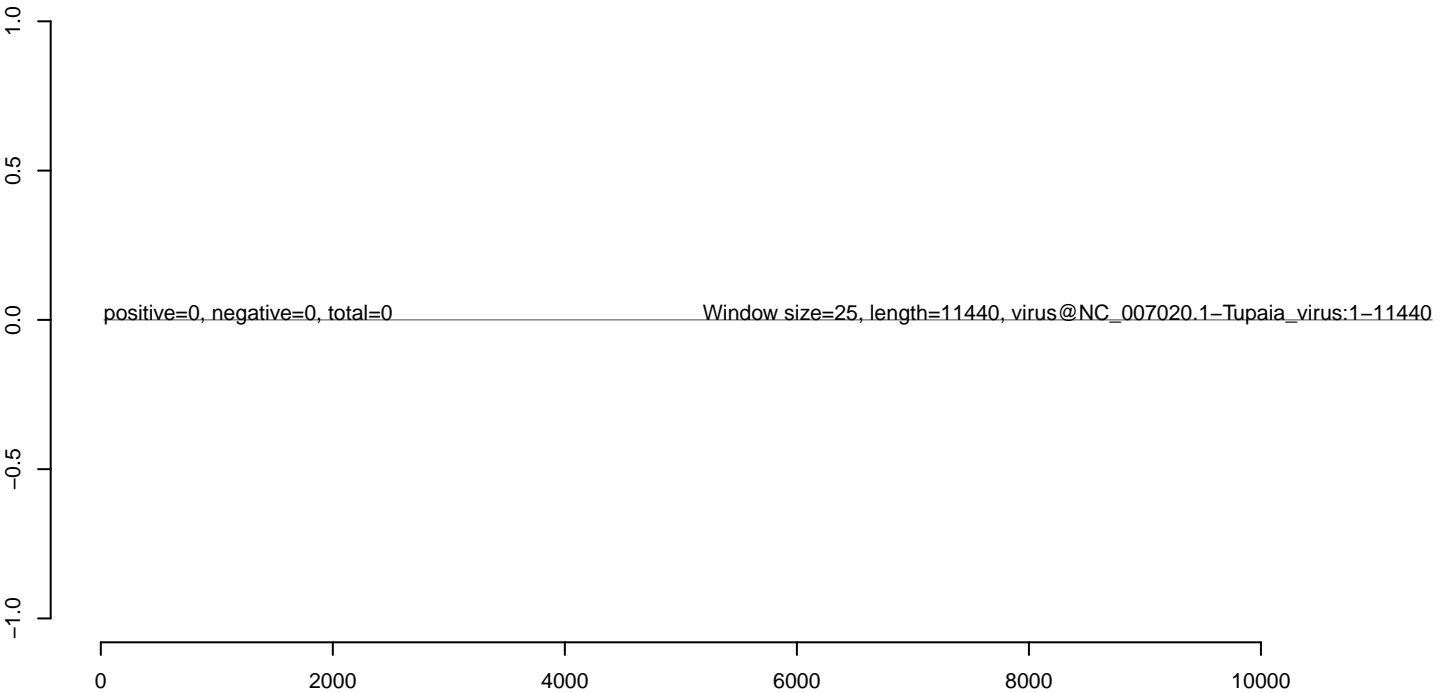
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



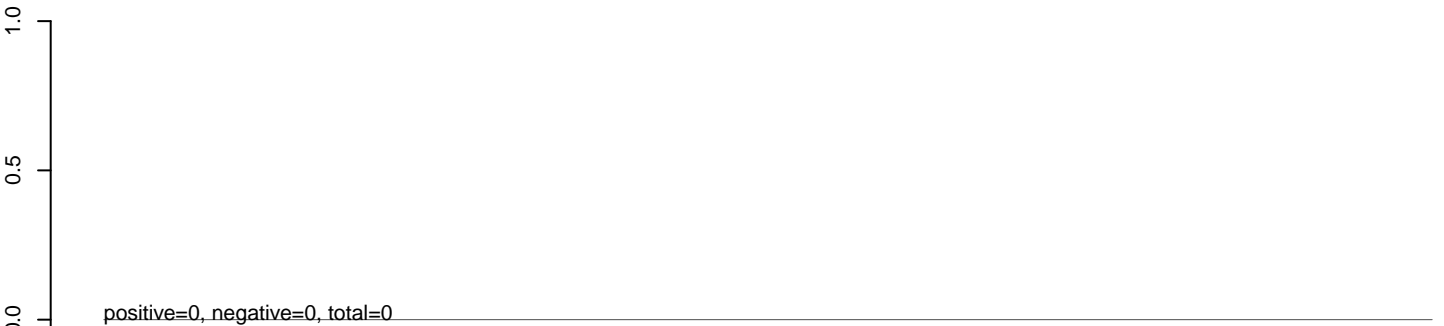
AeAlbo_C636_cells.rep



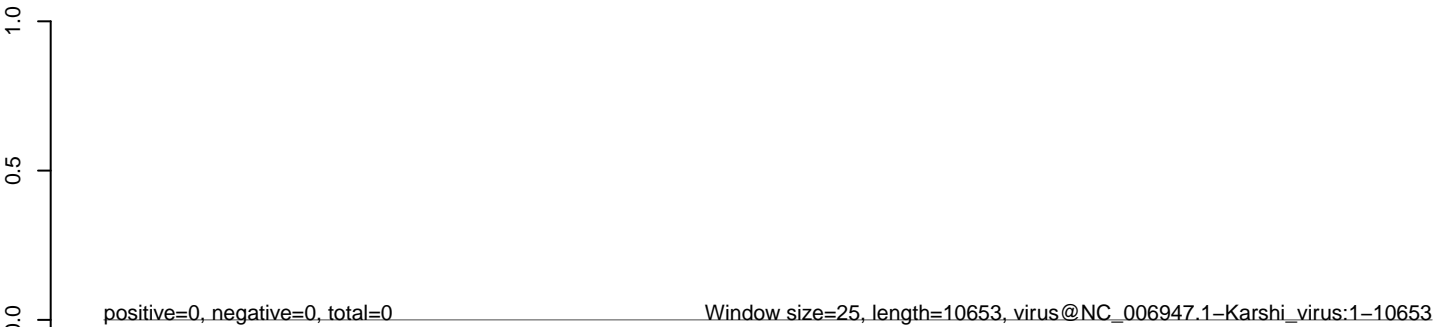
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

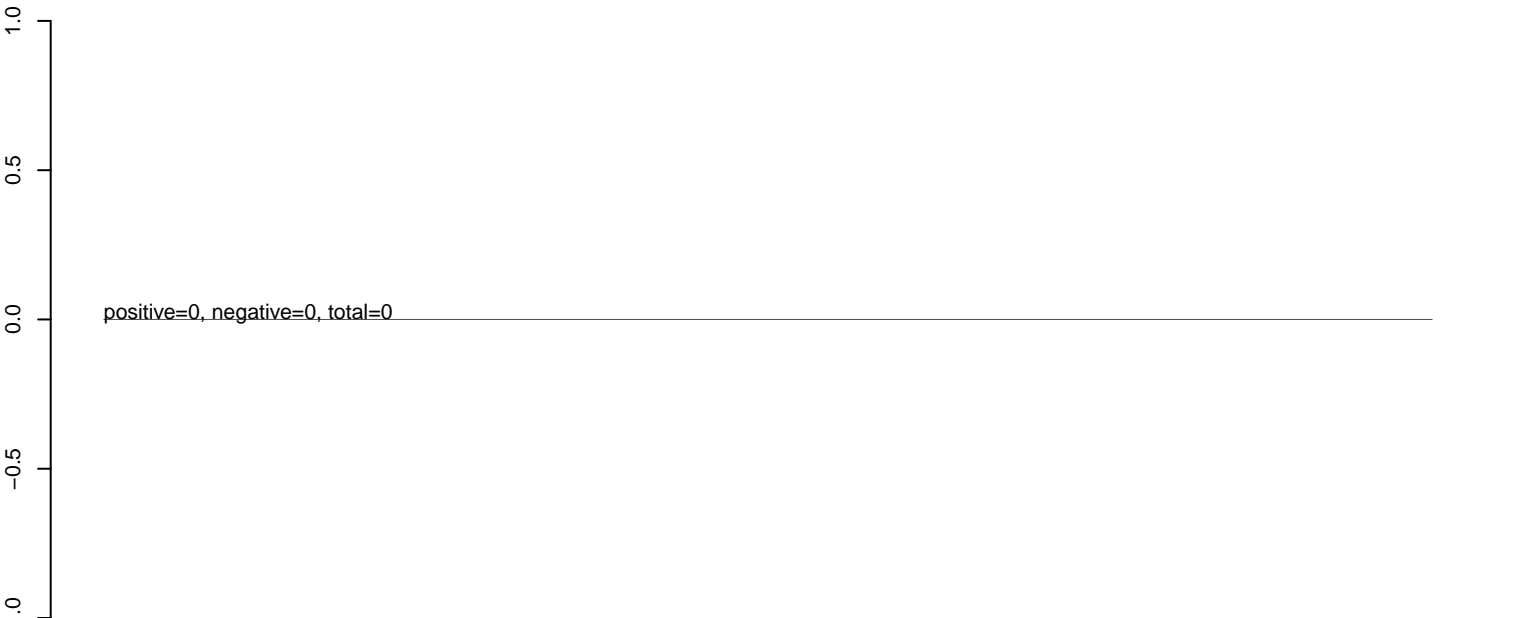


AeAlbo_C636_cells.rep

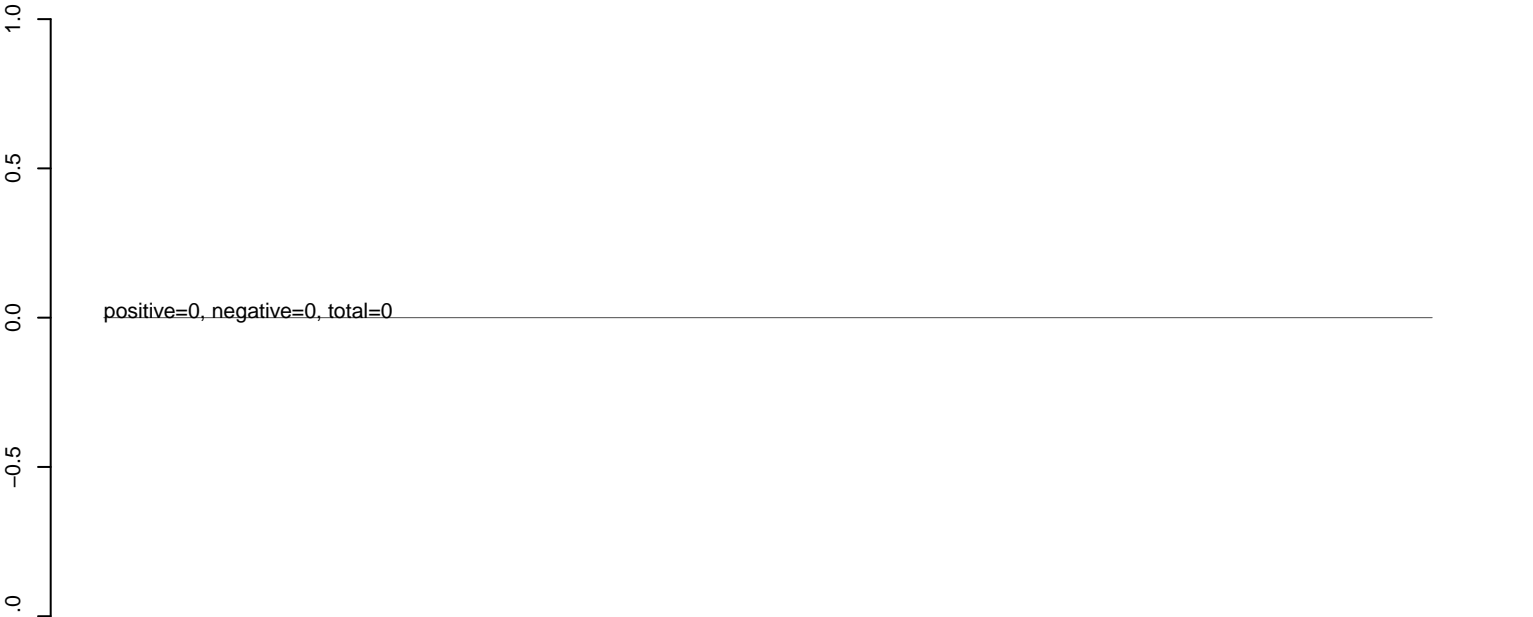


0 2000 4000 6000 8000 10000

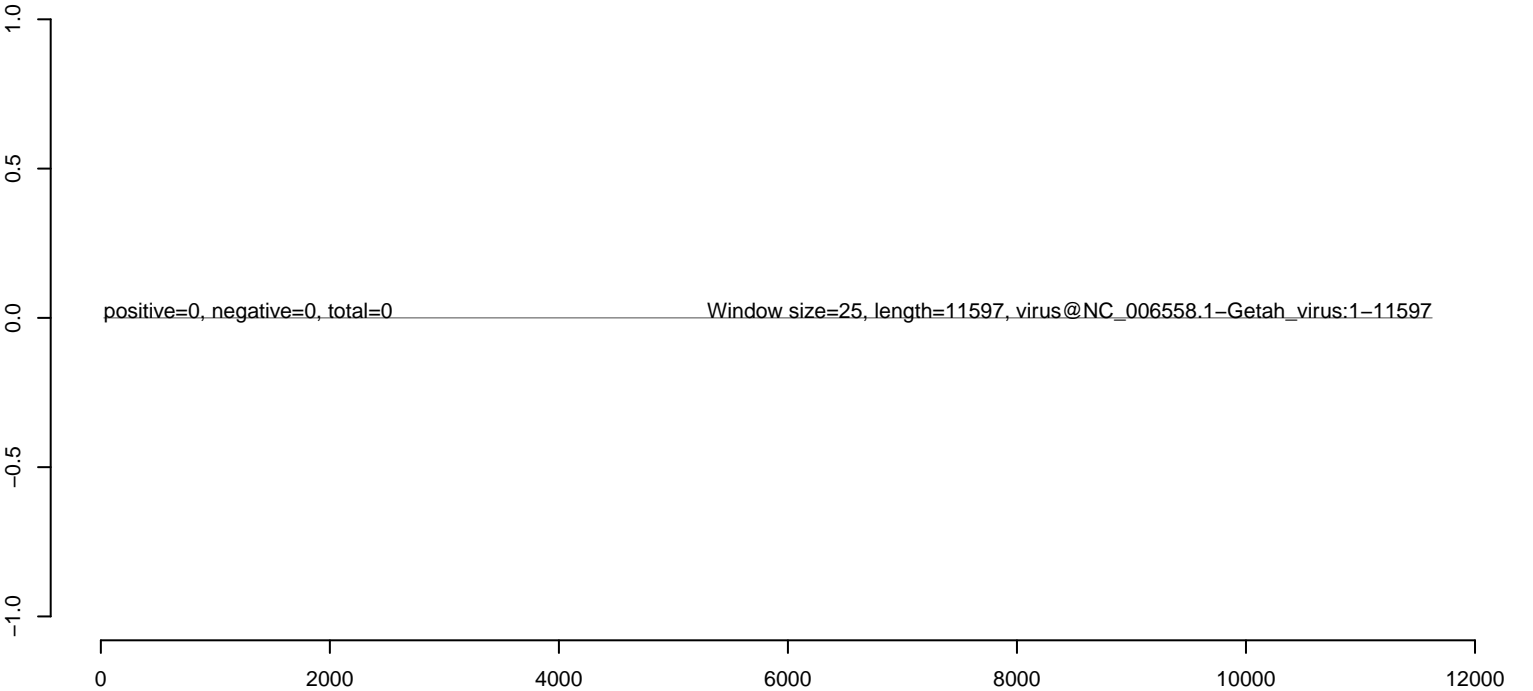
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



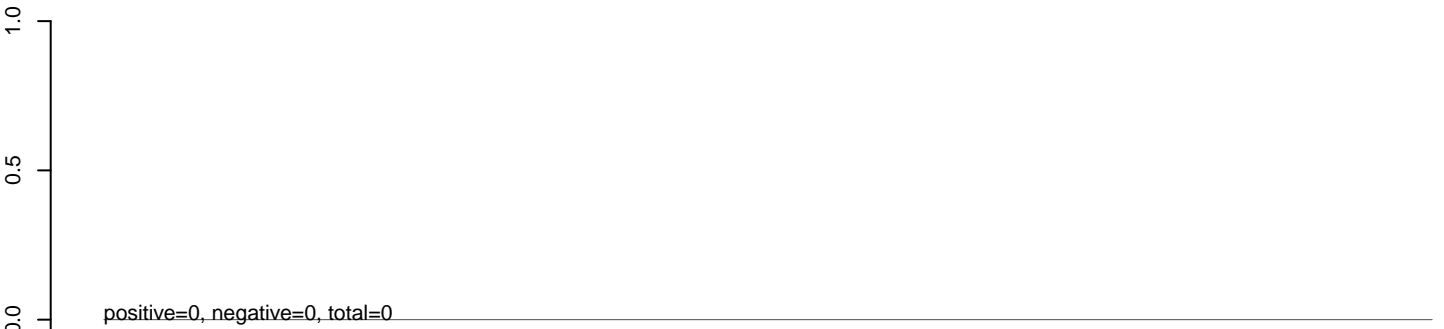
AeAlbo_C636_cells.rep



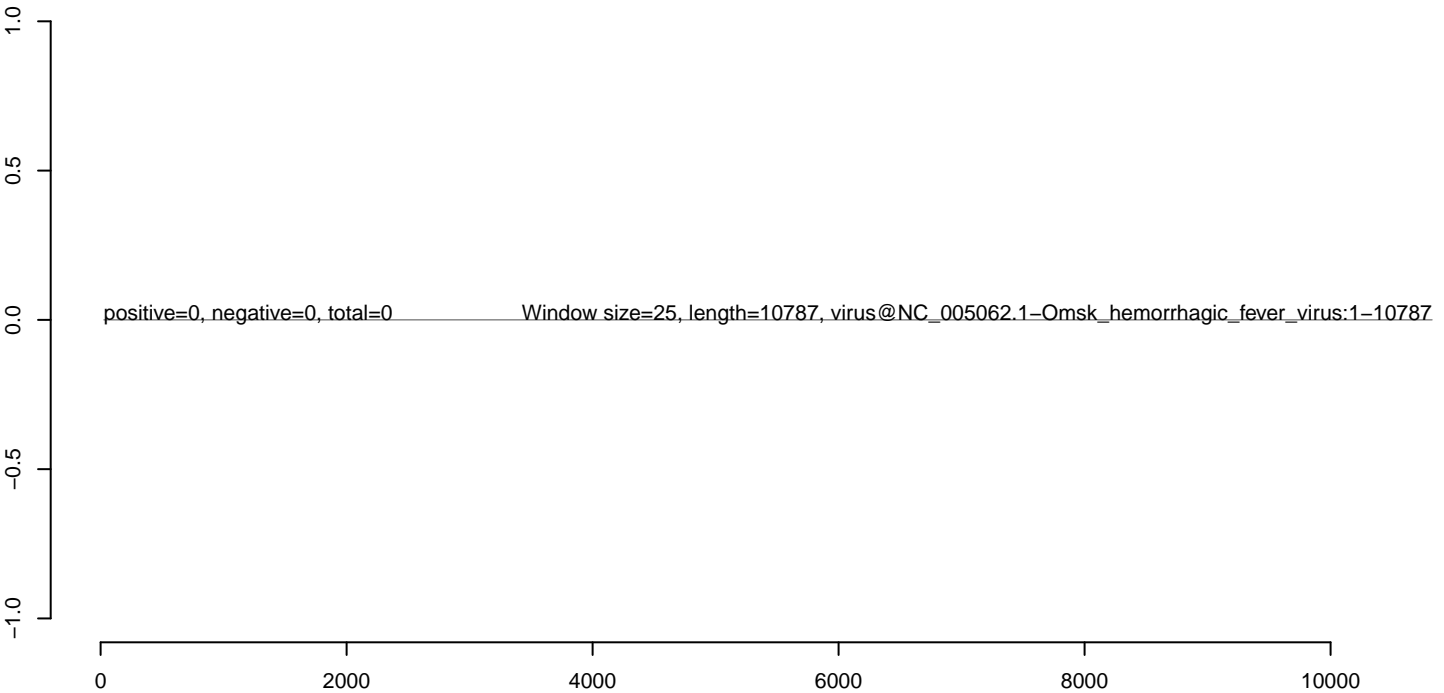
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



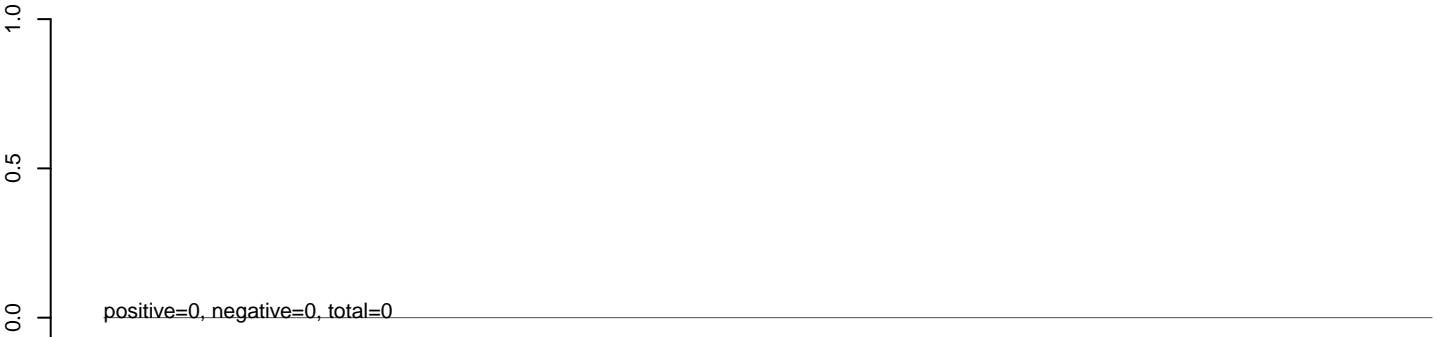
AeAlbo_C636_cells.rep



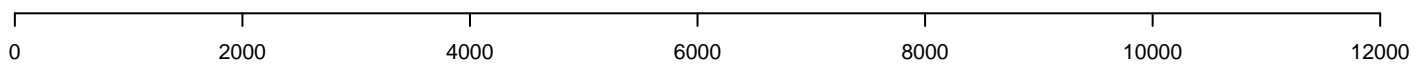
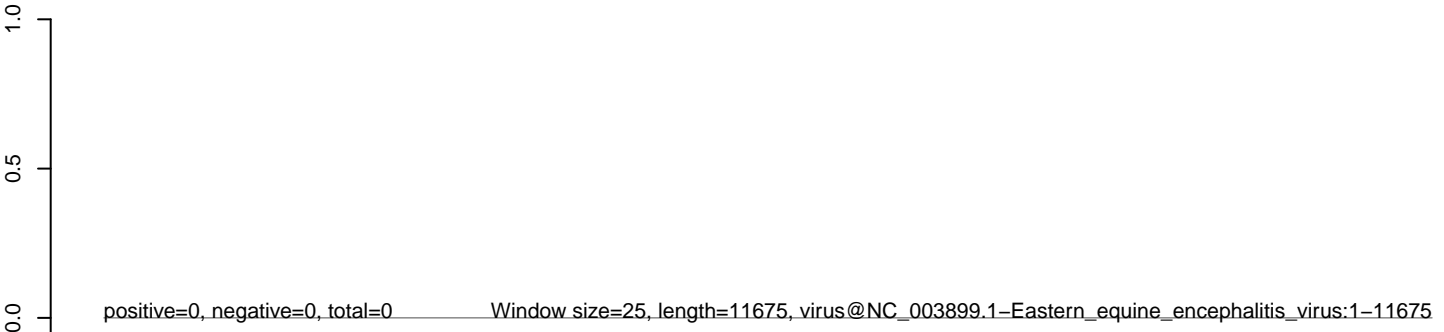
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



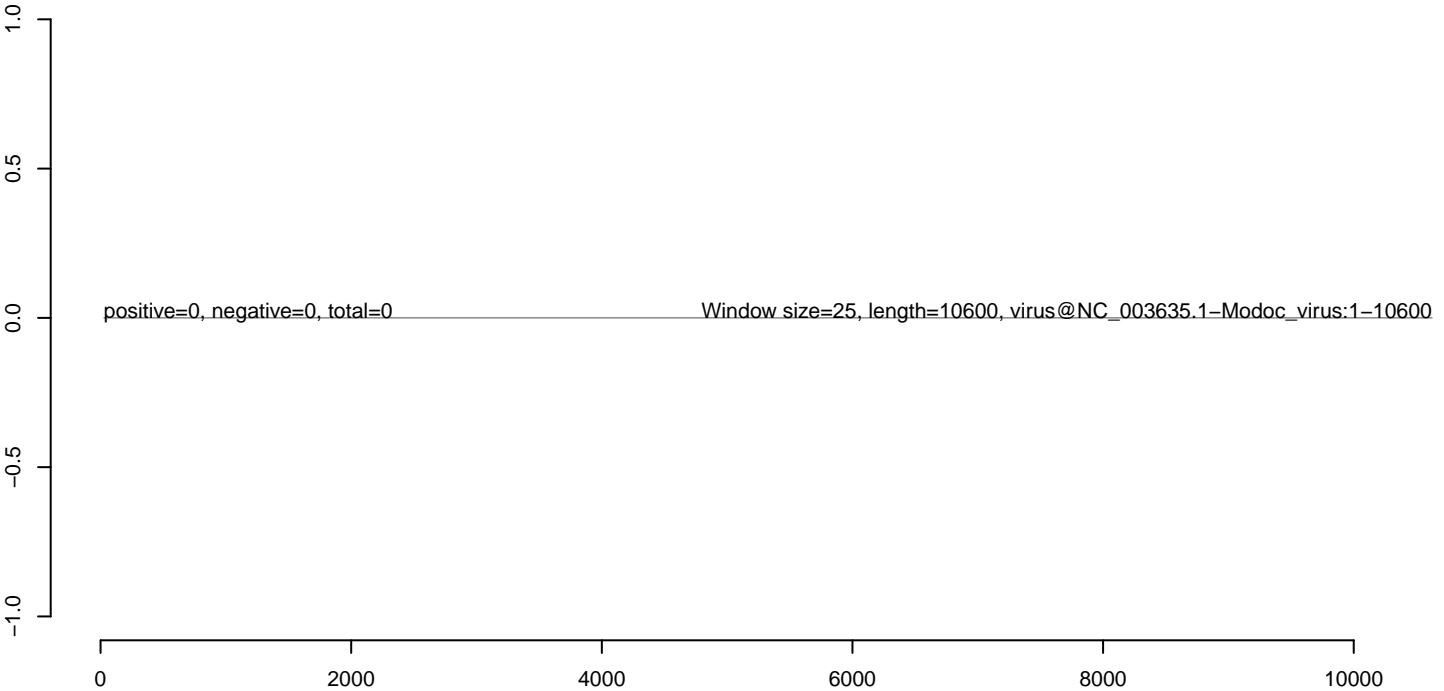
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



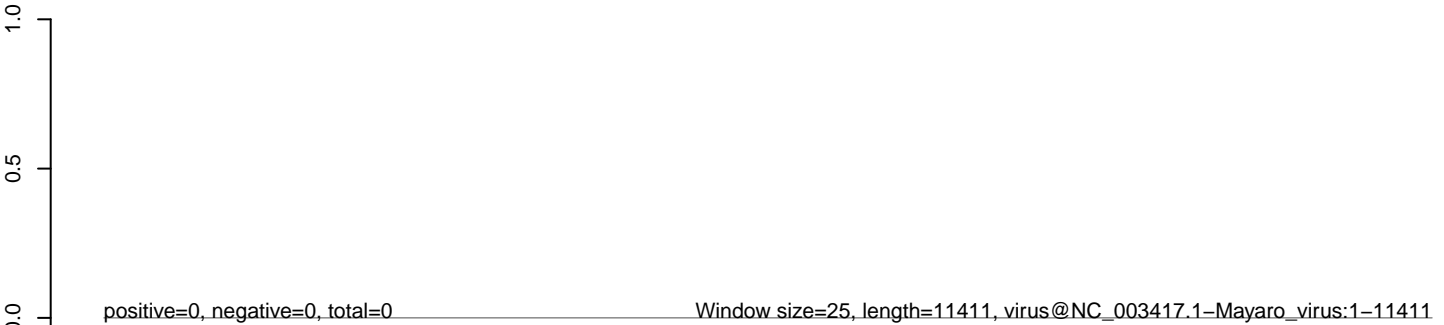
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

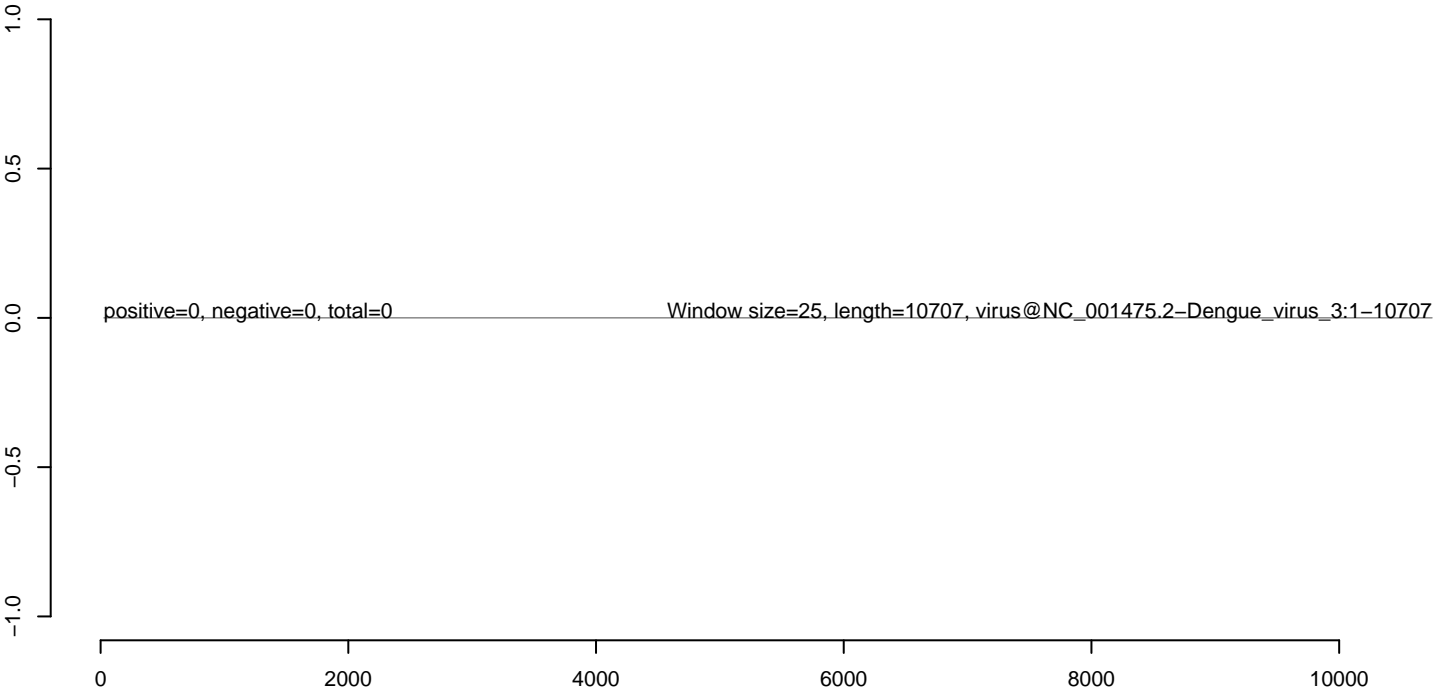
AeAlbo_C636_cells.18_23.rep



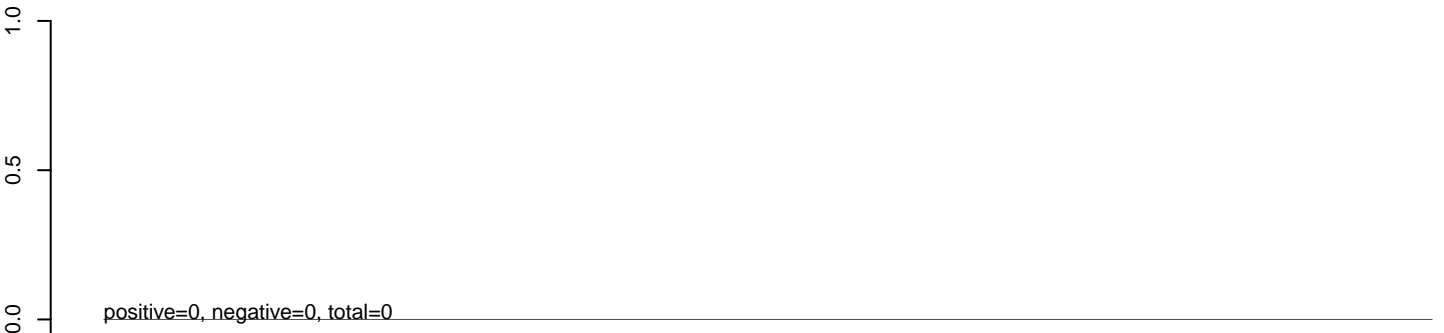
AeAlbo_C636_cells.24_35.rep



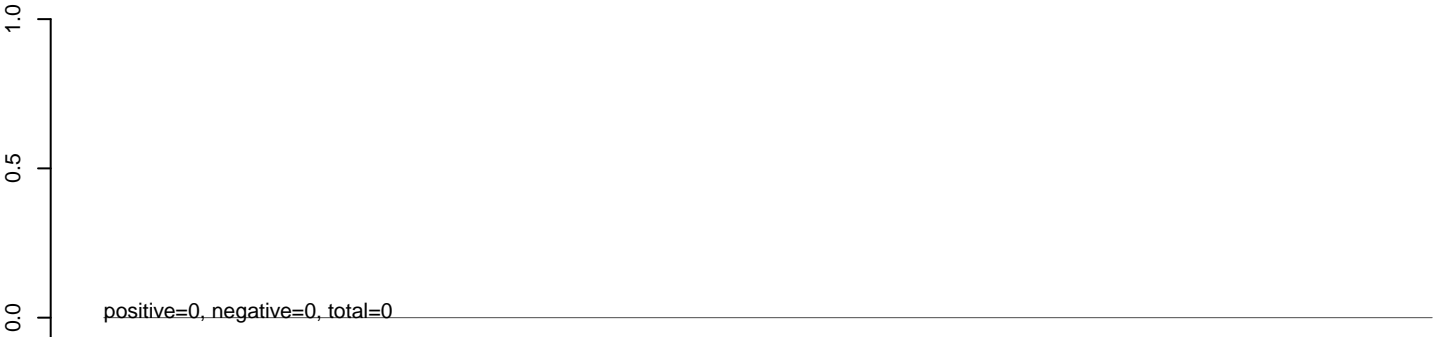
AeAlbo_C636_cells.rep



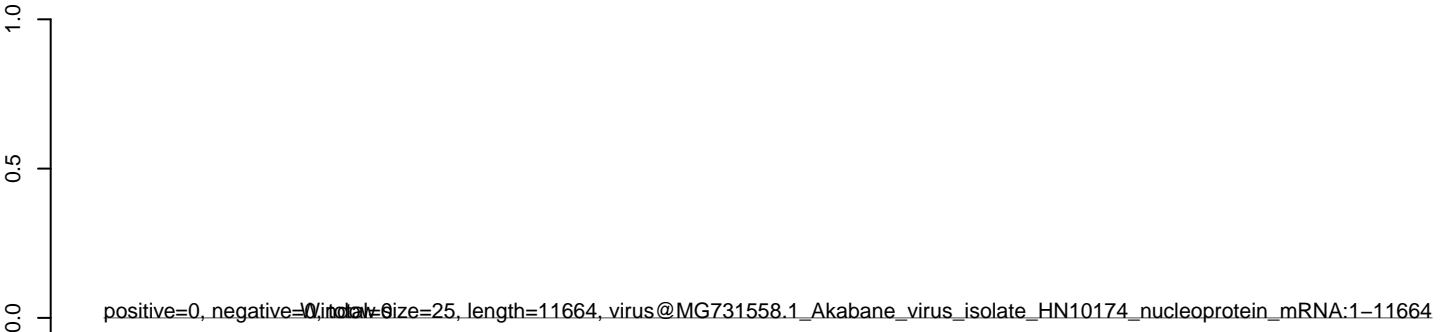
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000 12000

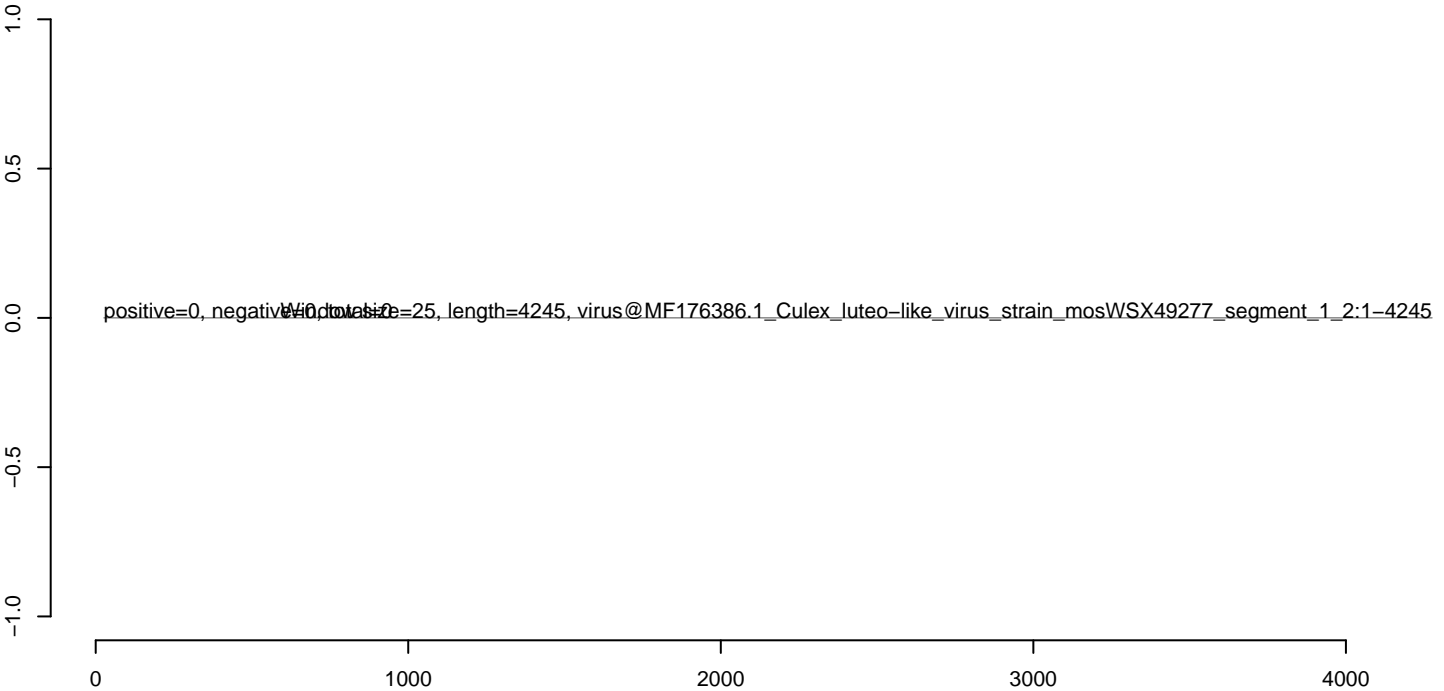
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



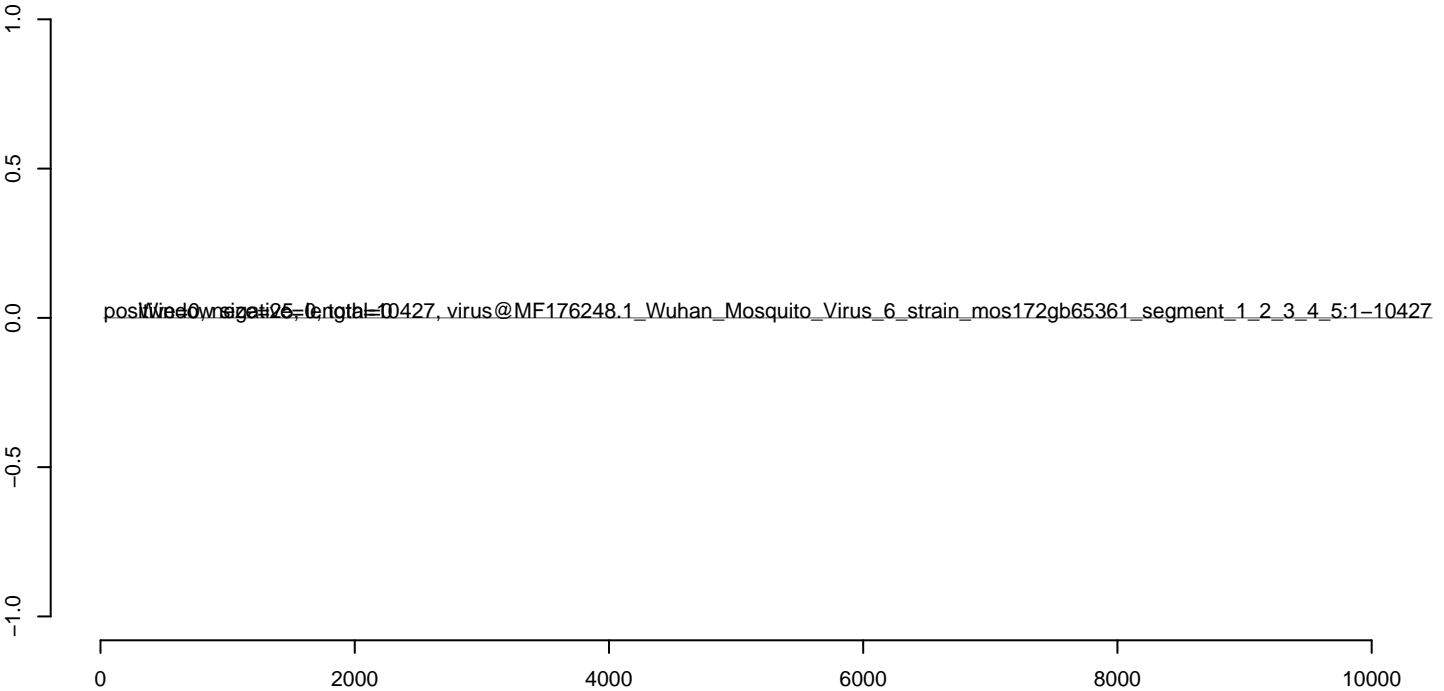
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



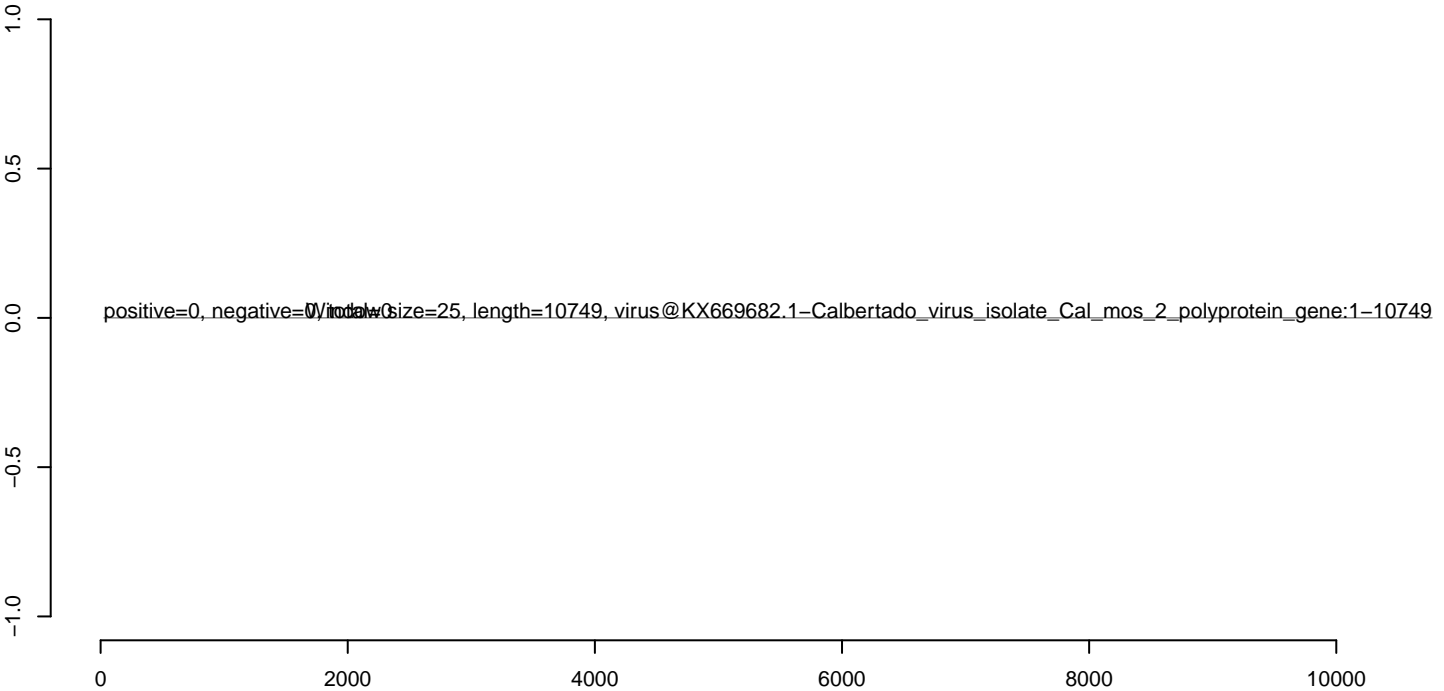
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



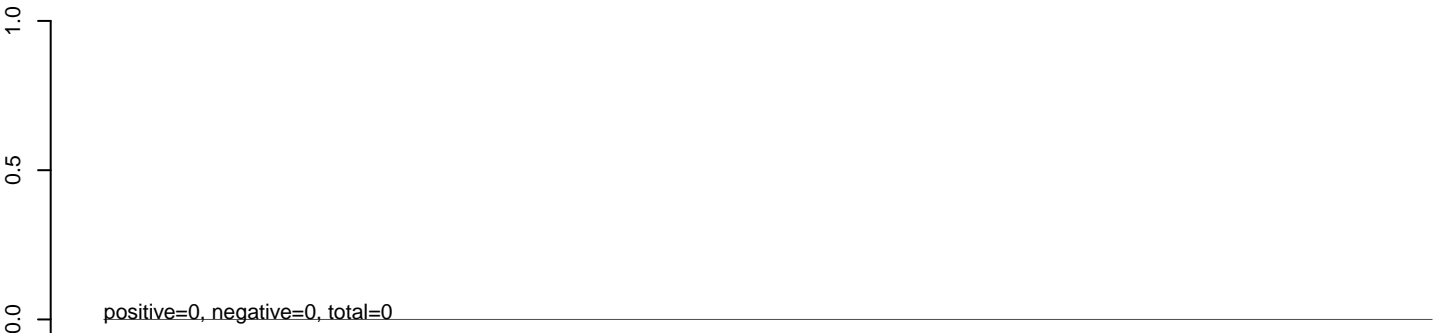
AeAlbo_C636_cells.24_35.rep



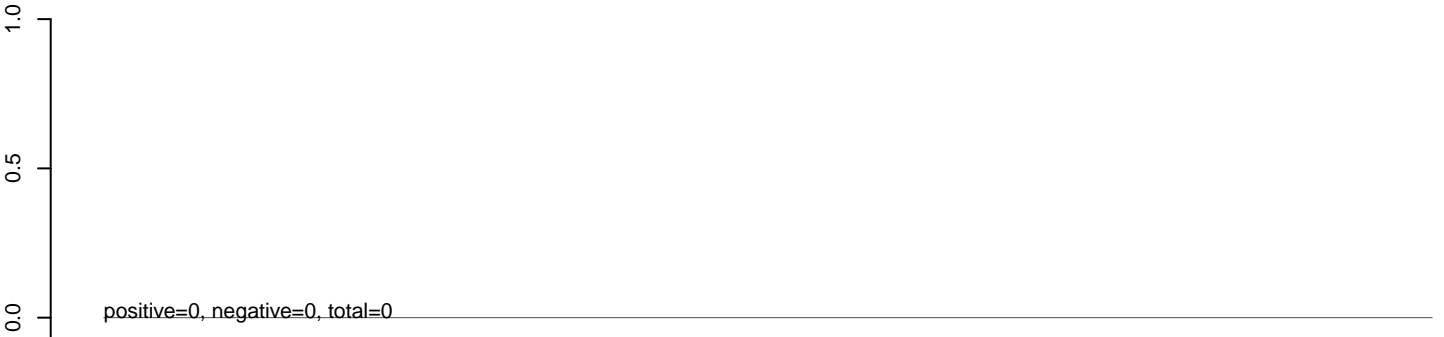
AeAlbo_C636_cells.rep



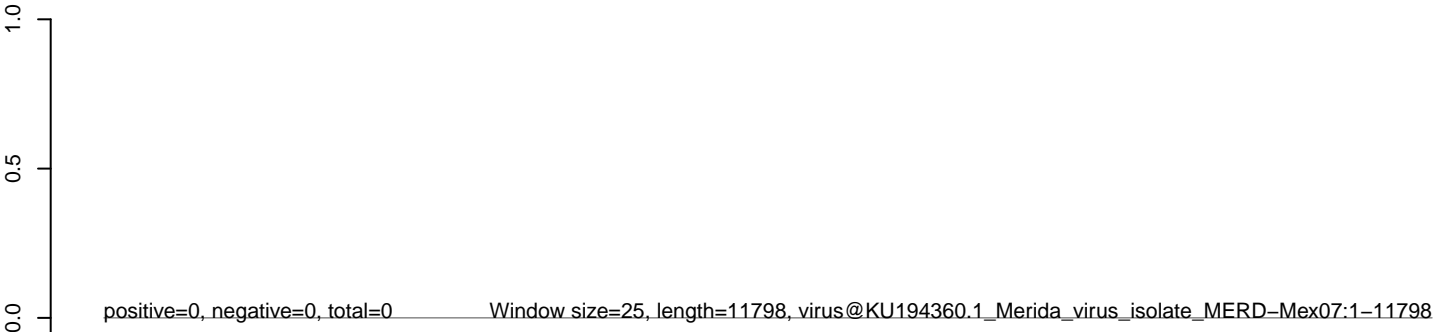
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep

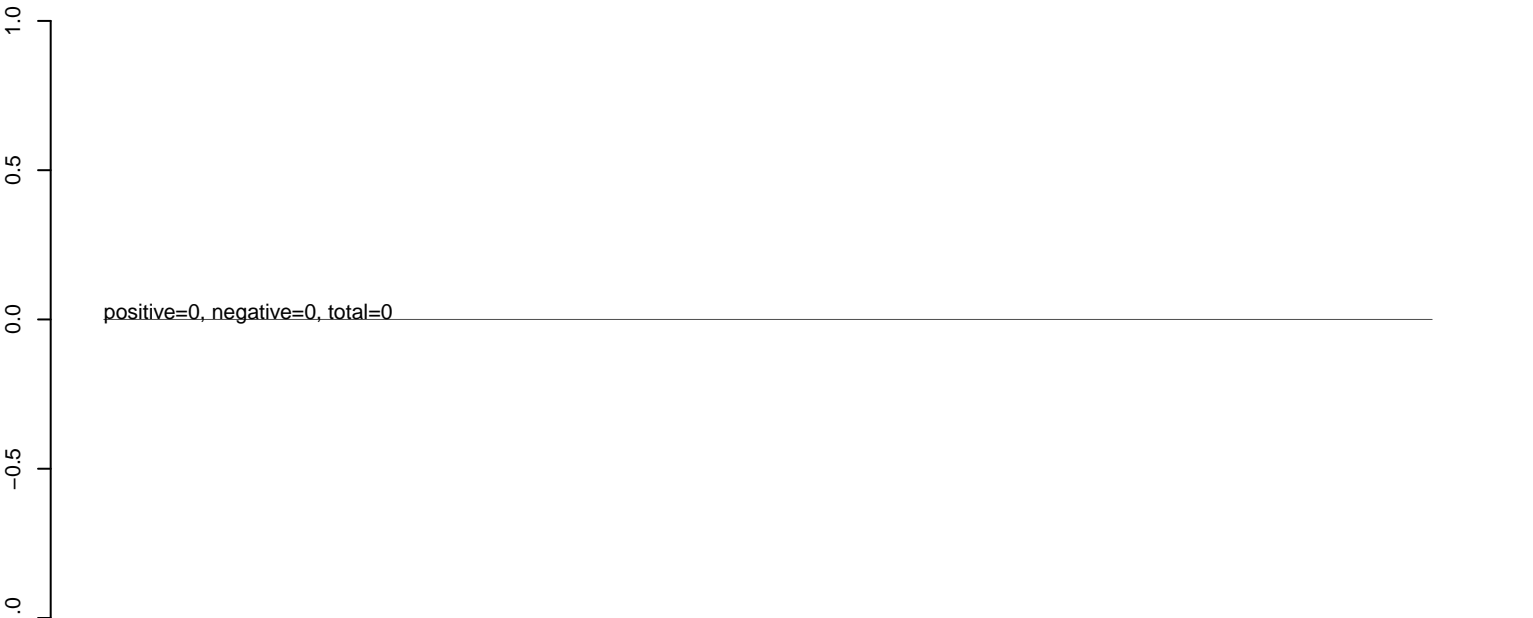


AeAlbo_C636_cells.rep

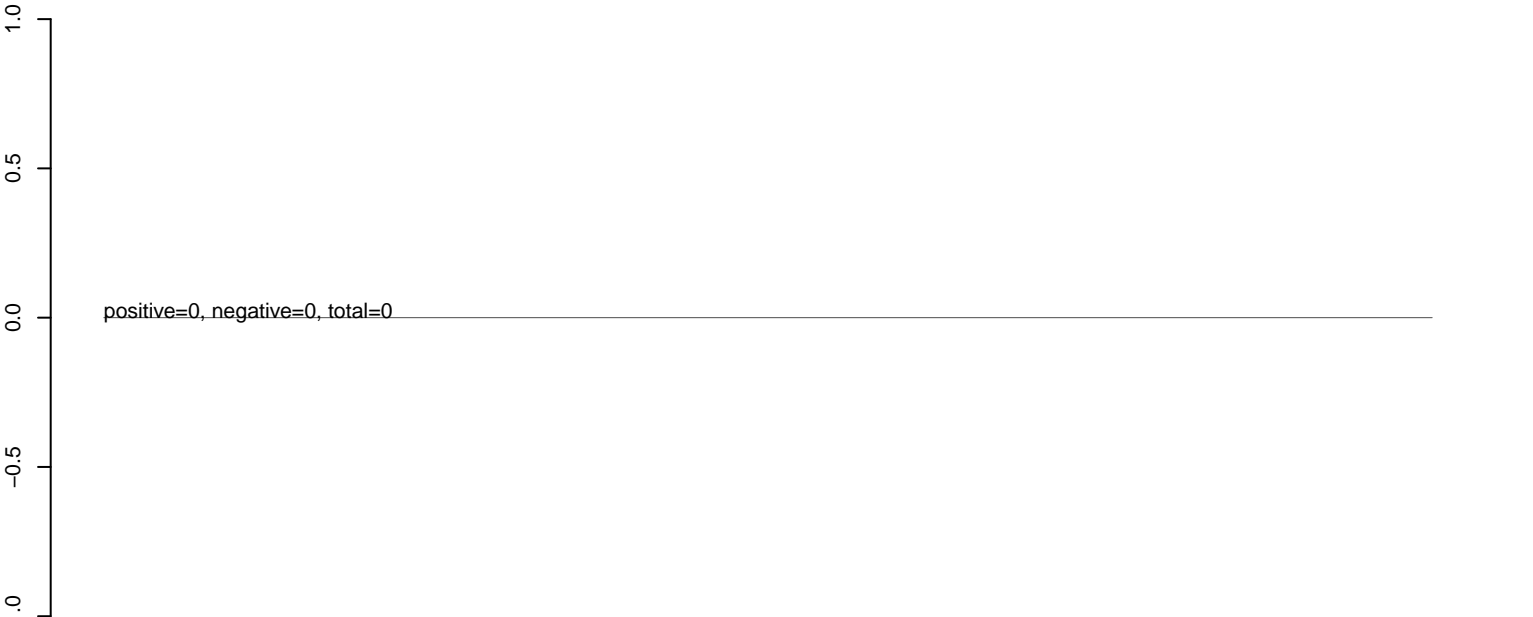


0 2000 4000 6000 8000 10000 12000

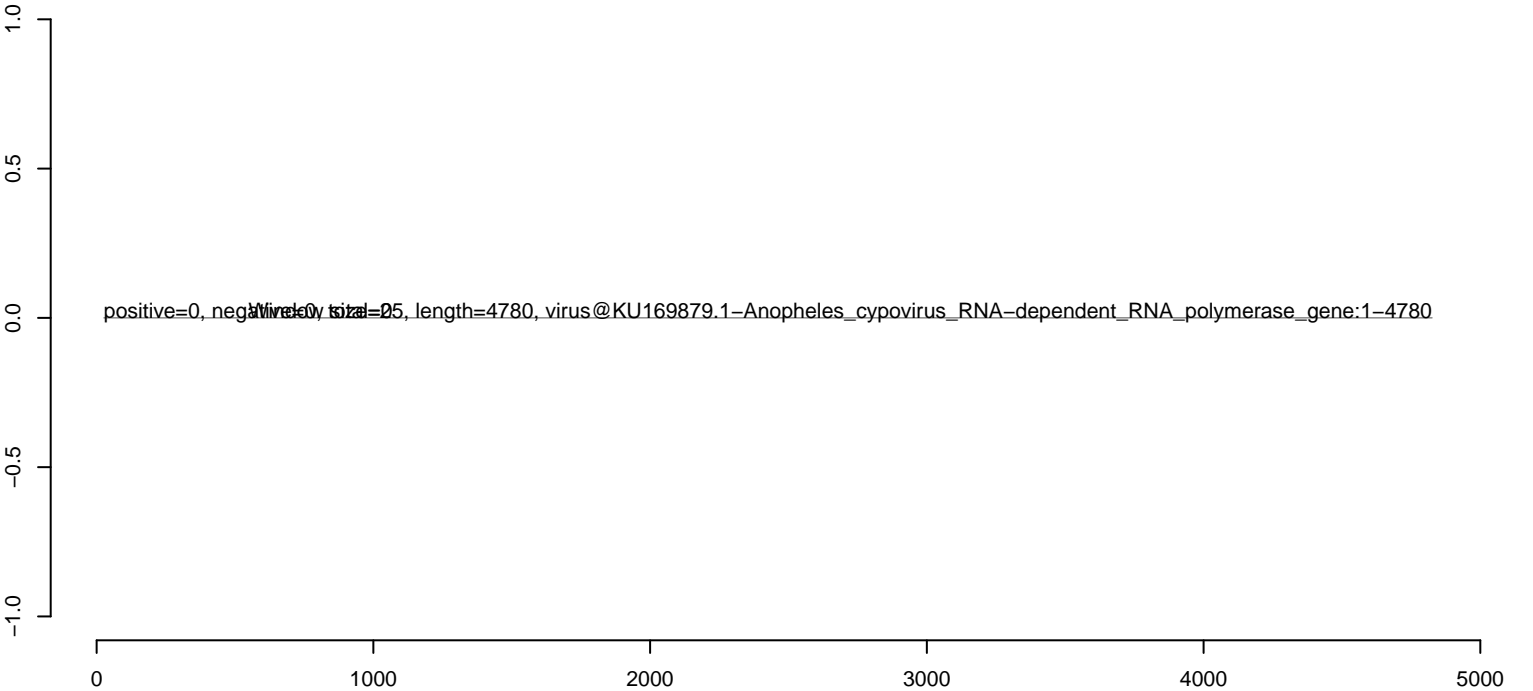
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



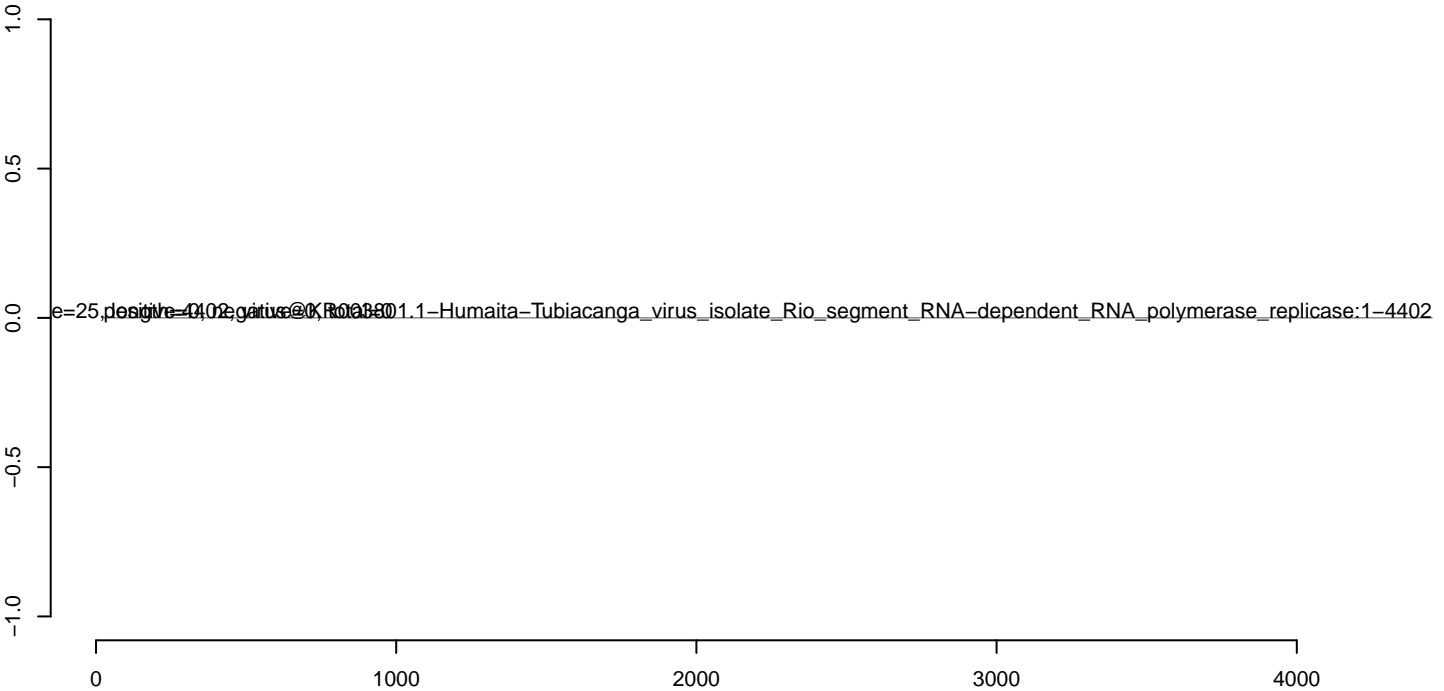
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



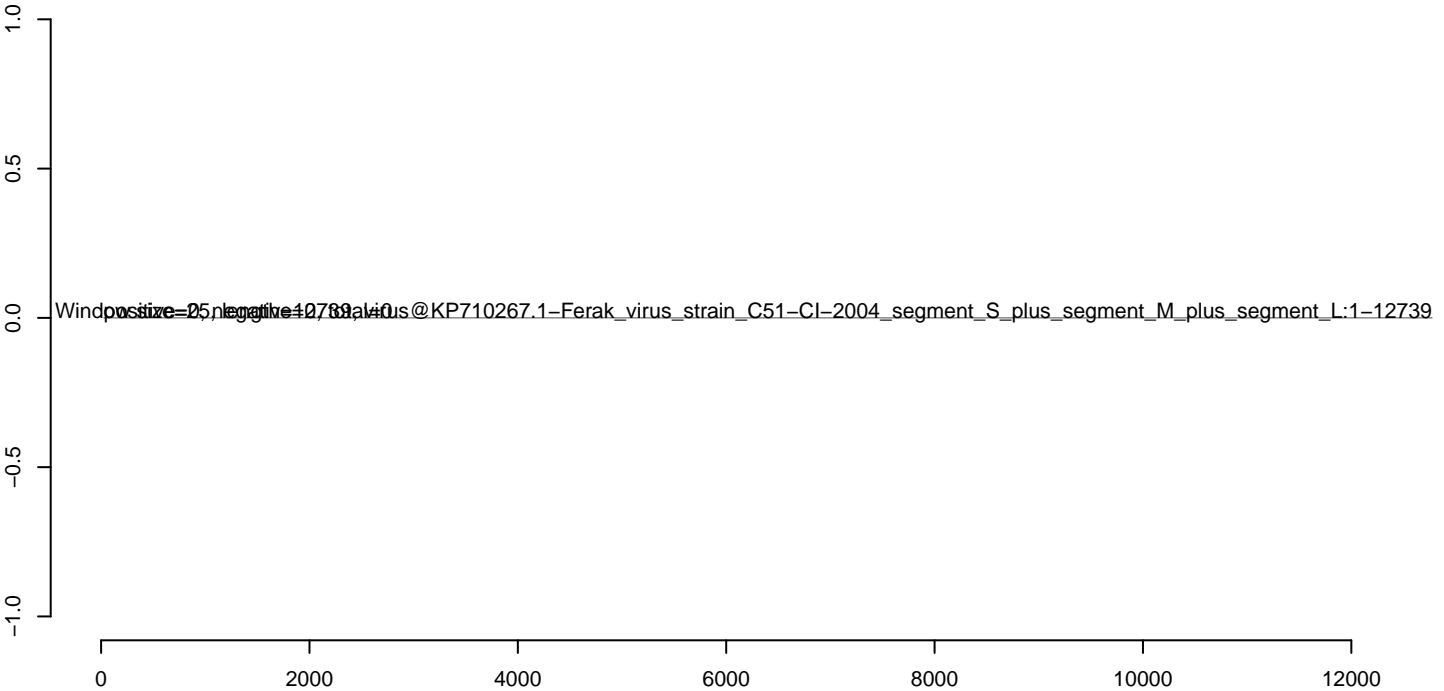
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



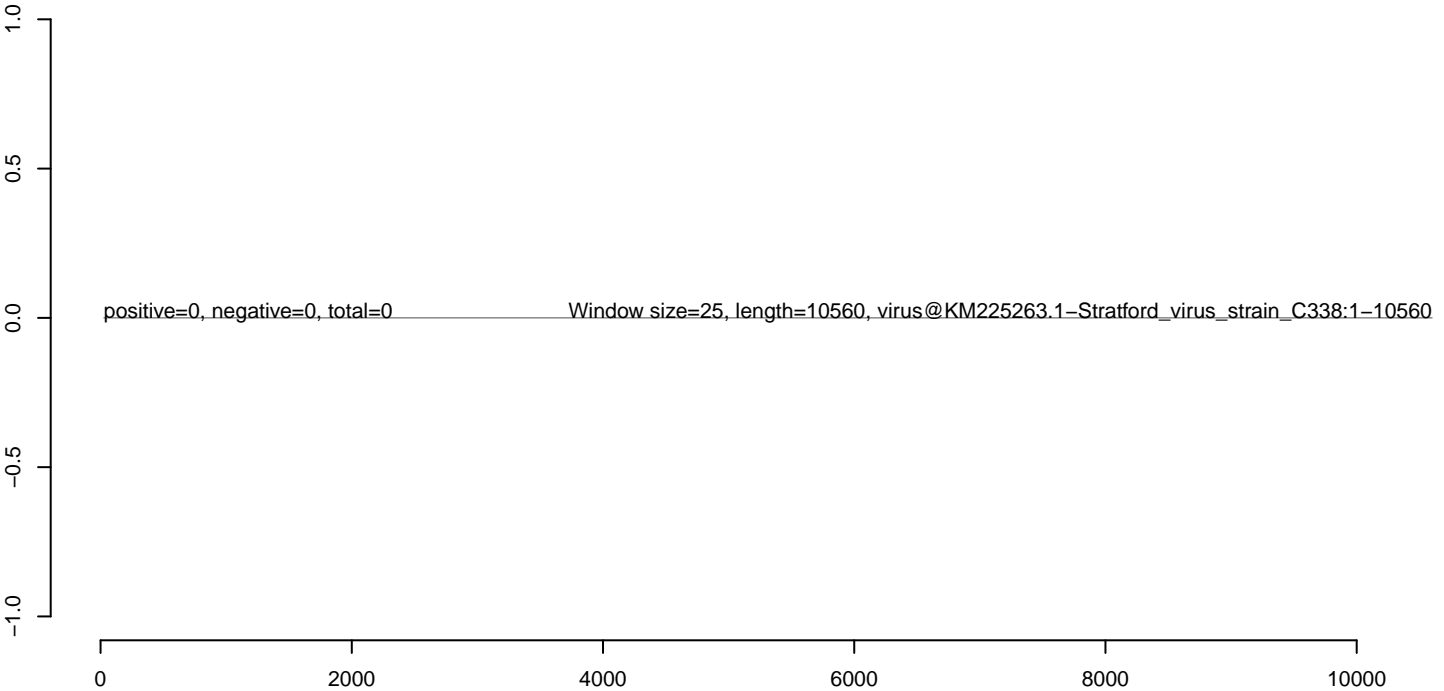
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



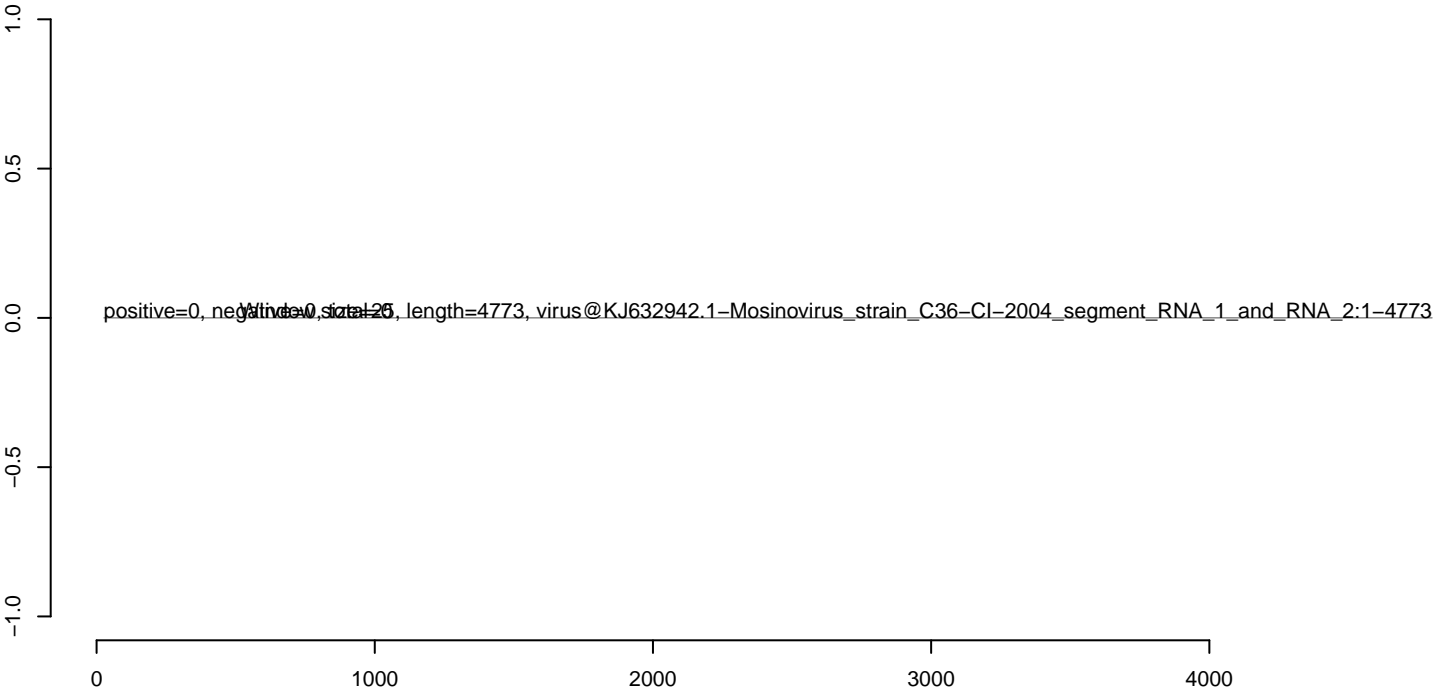
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



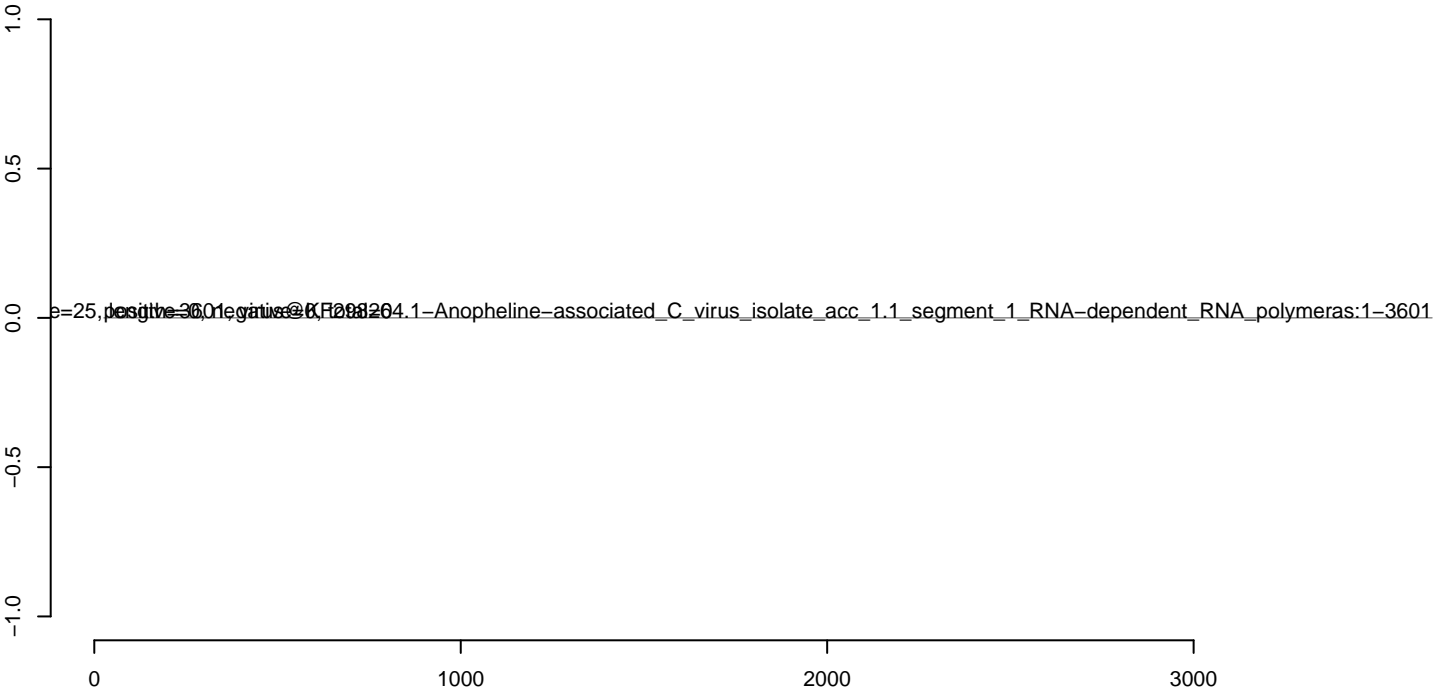
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



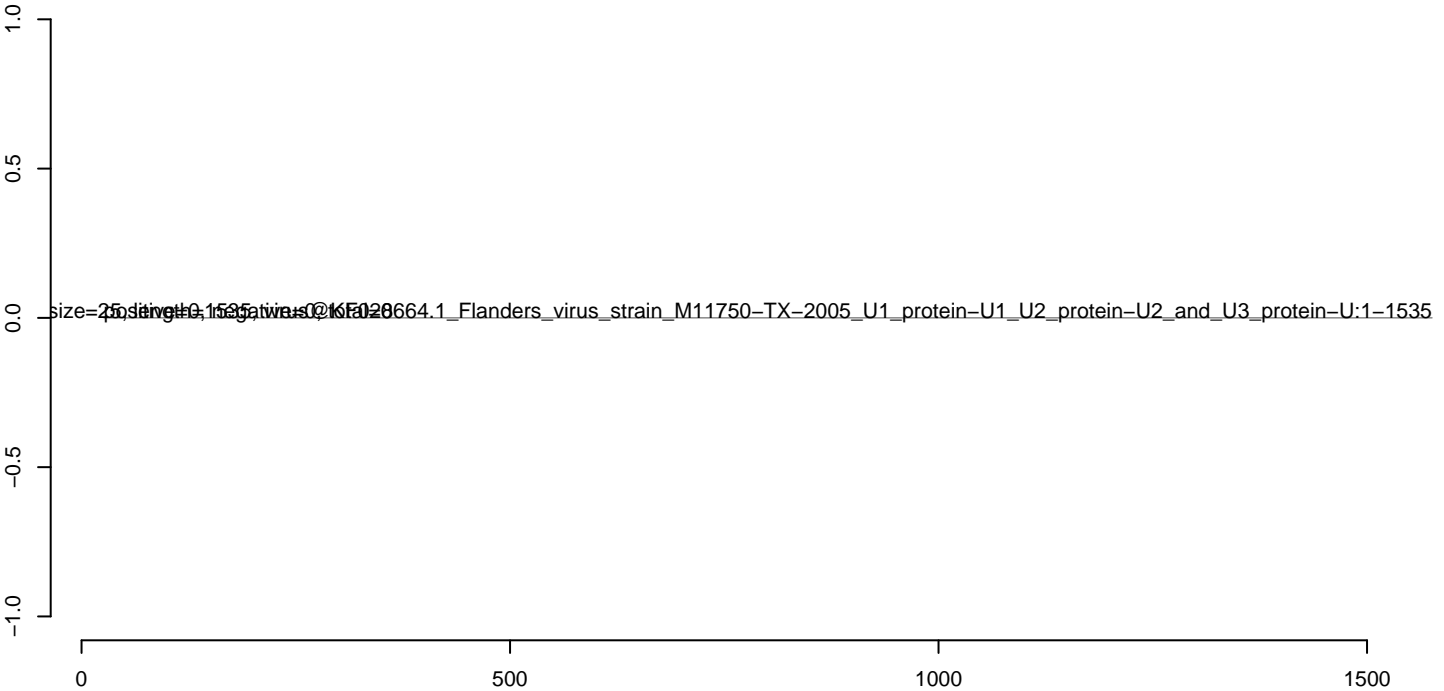
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



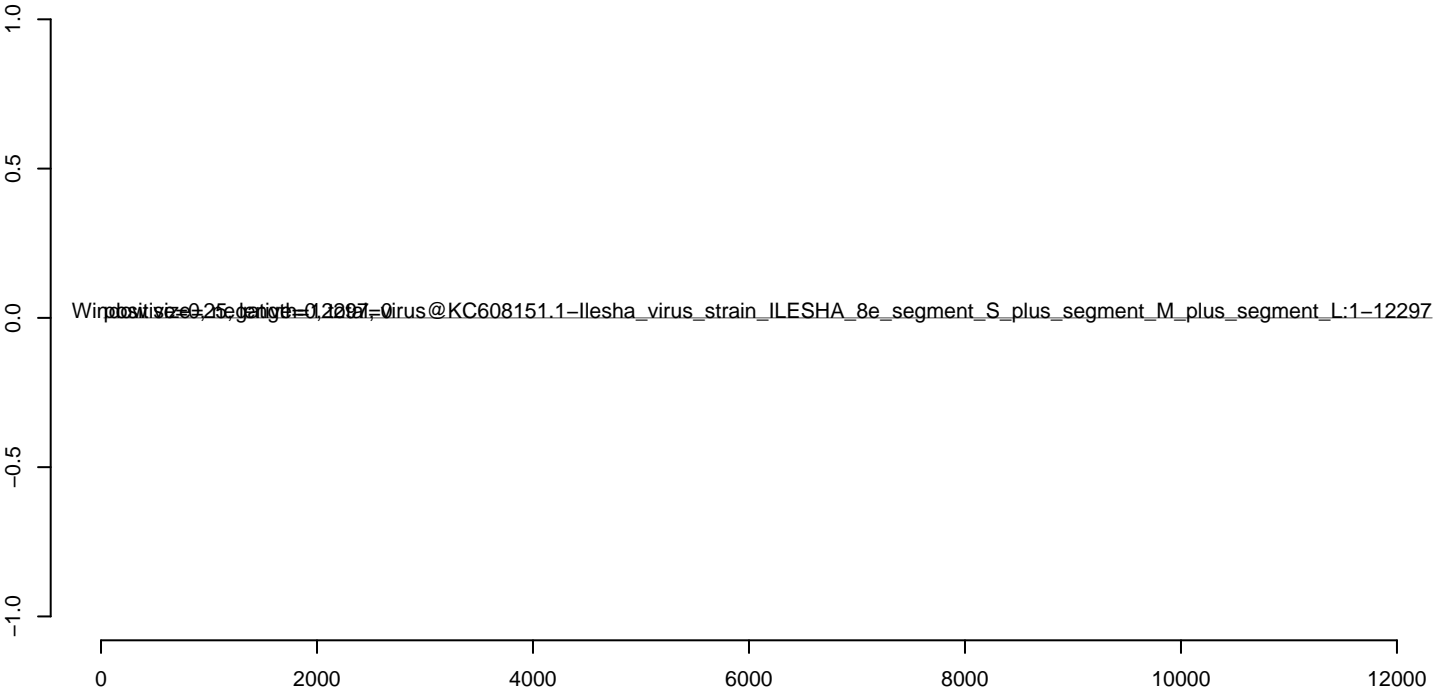
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



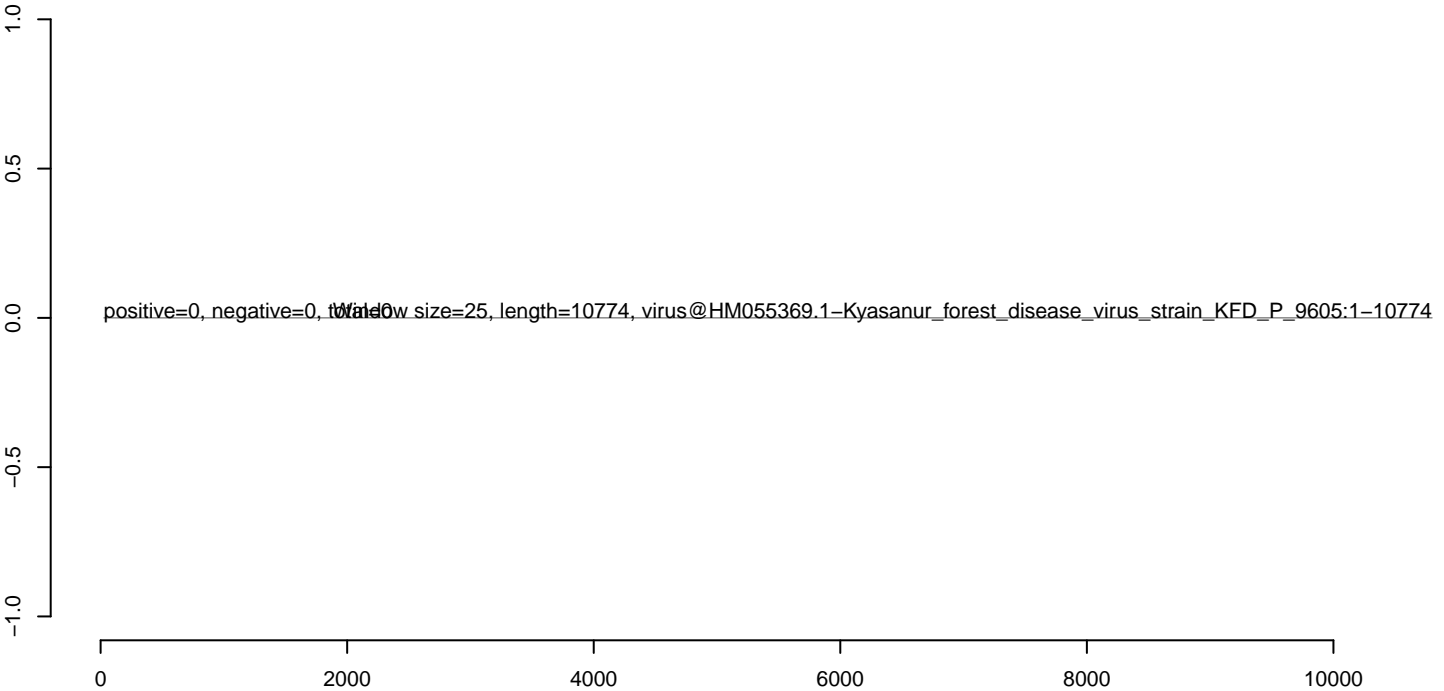
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



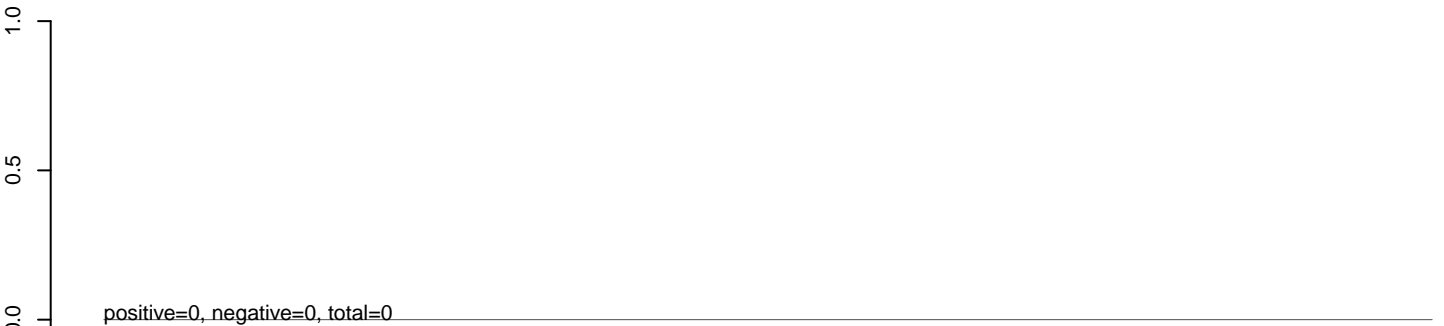
AeAlbo_C636_cells.rep



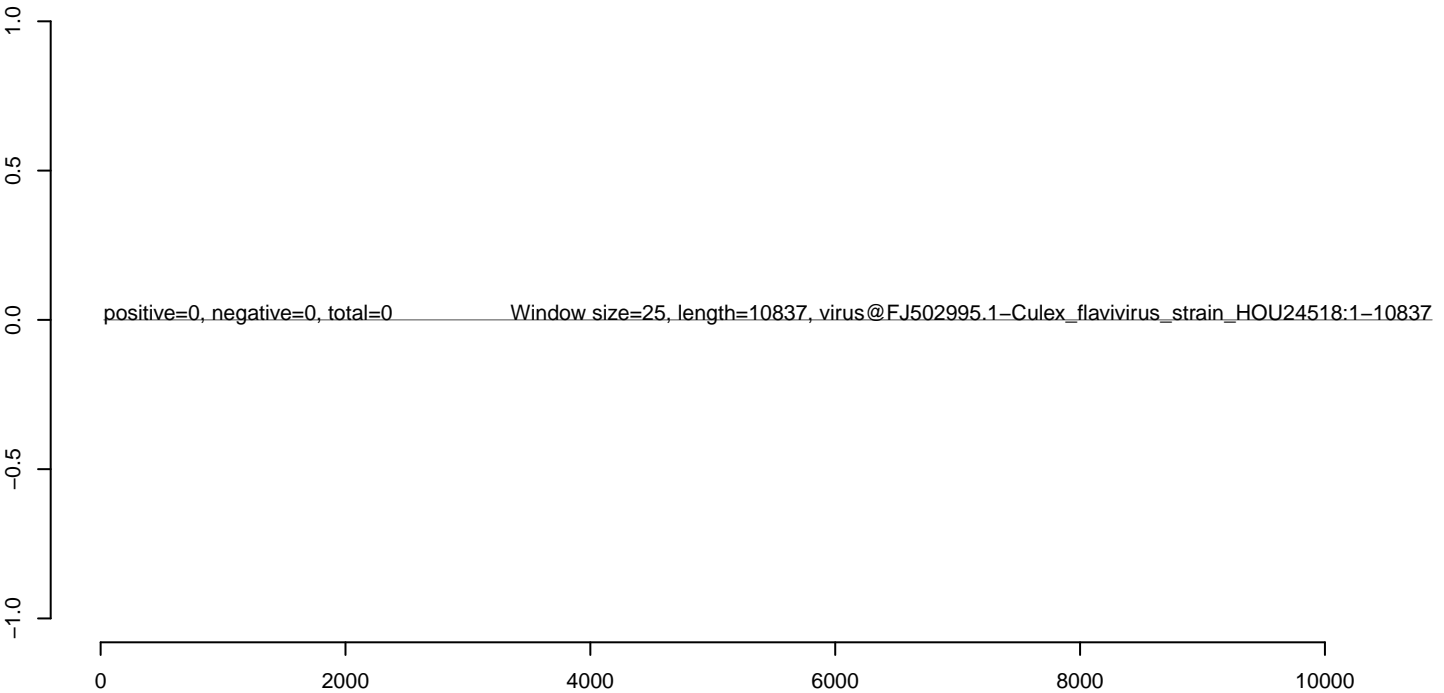
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



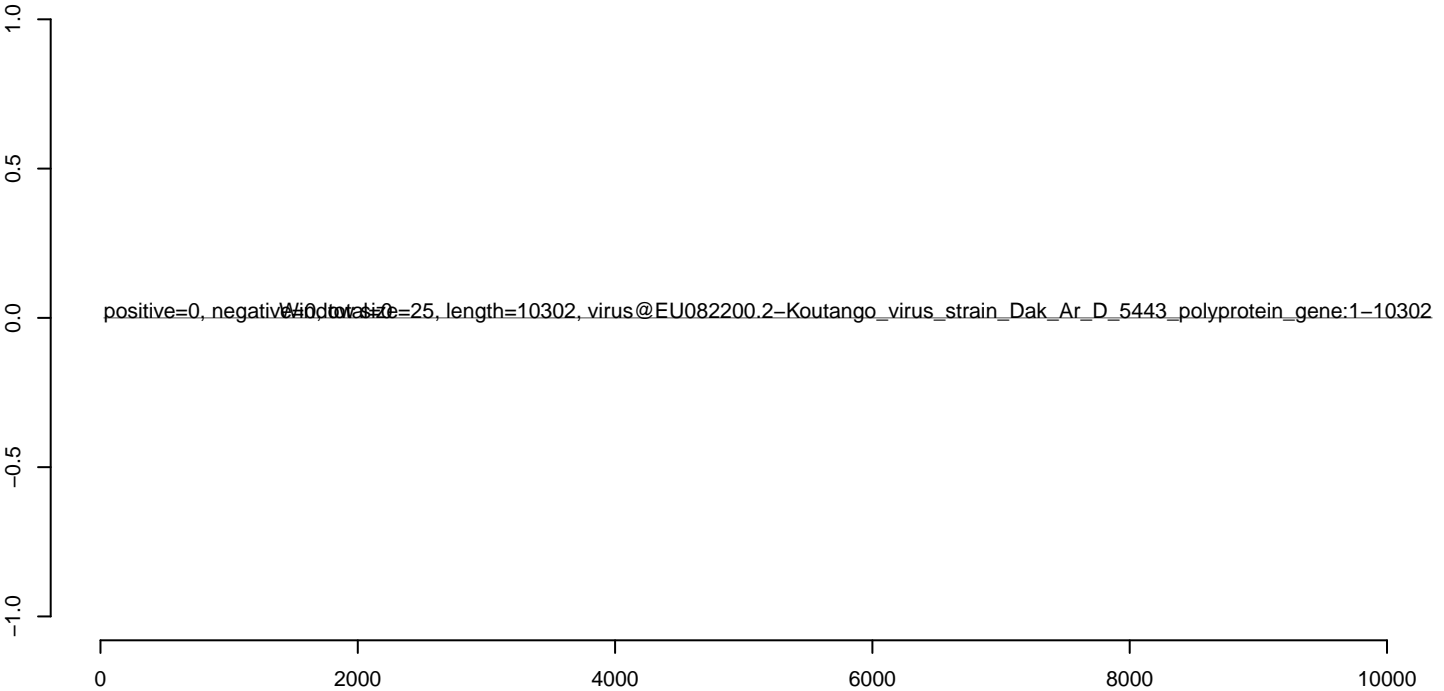
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



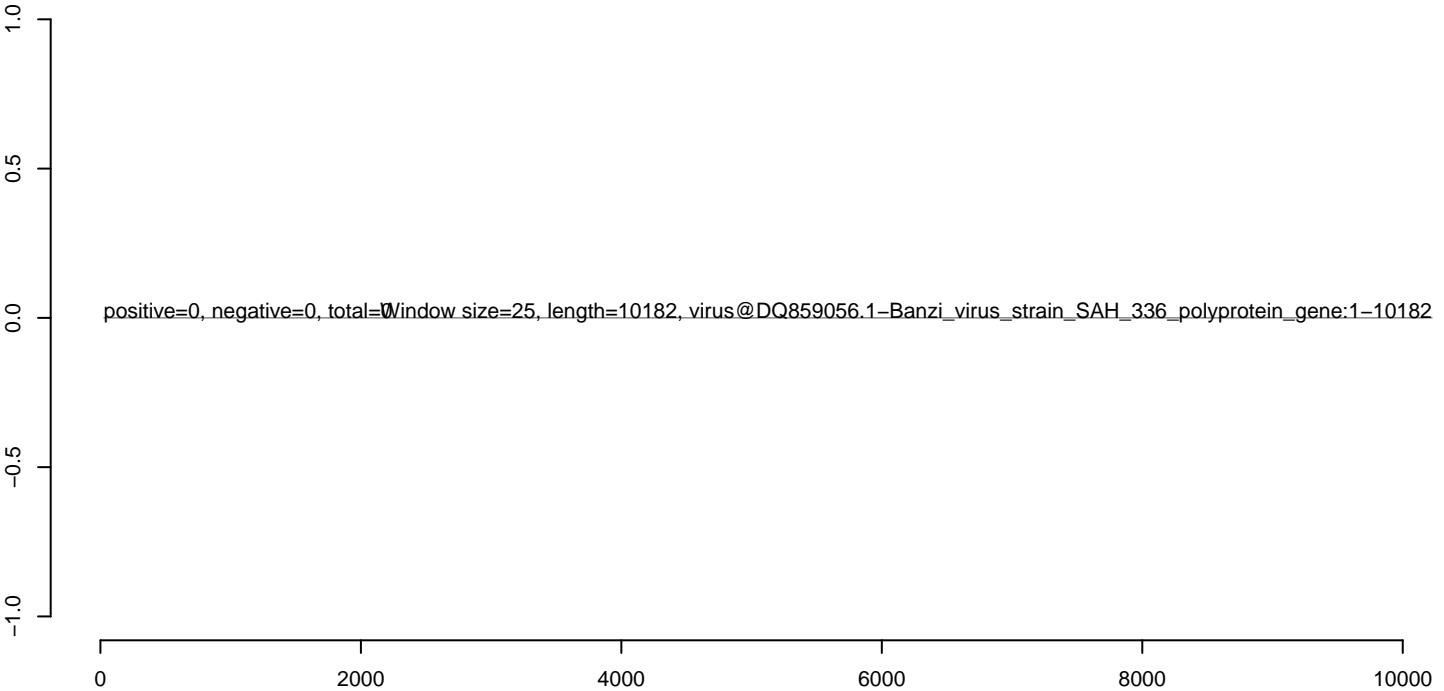
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



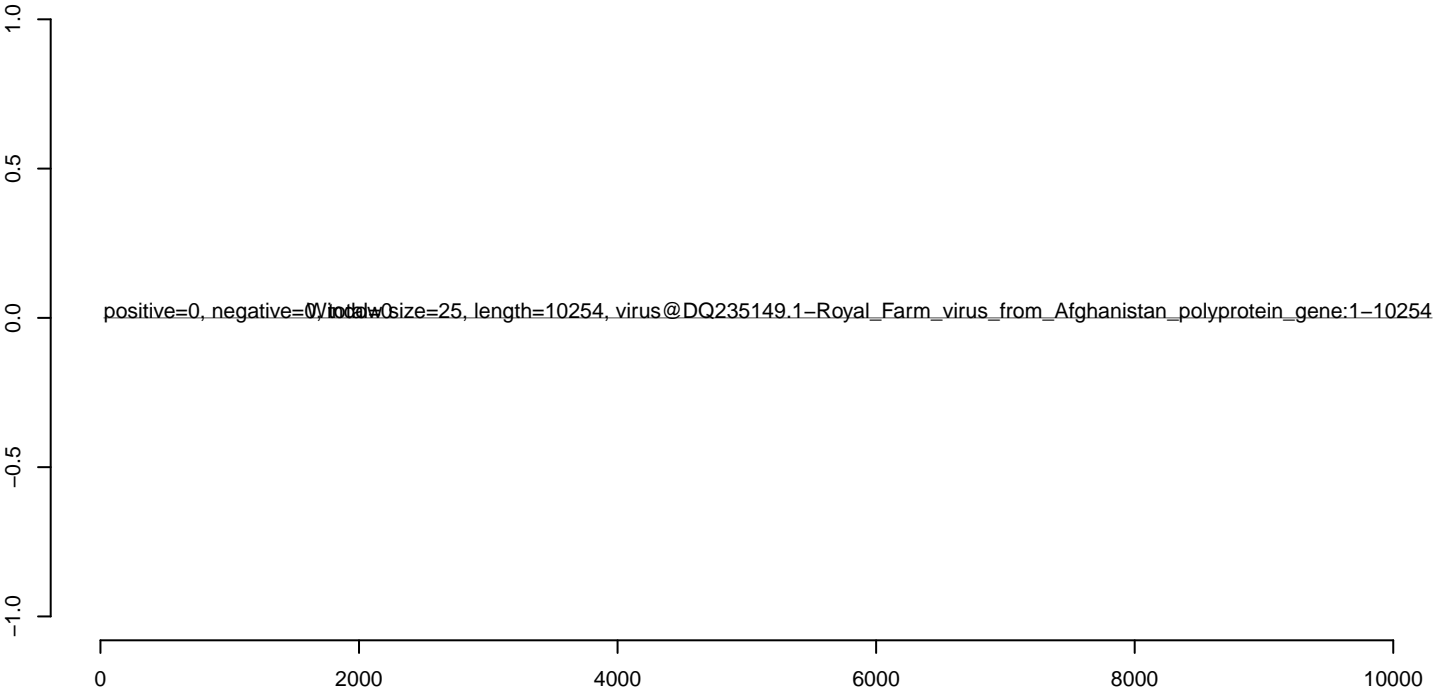
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



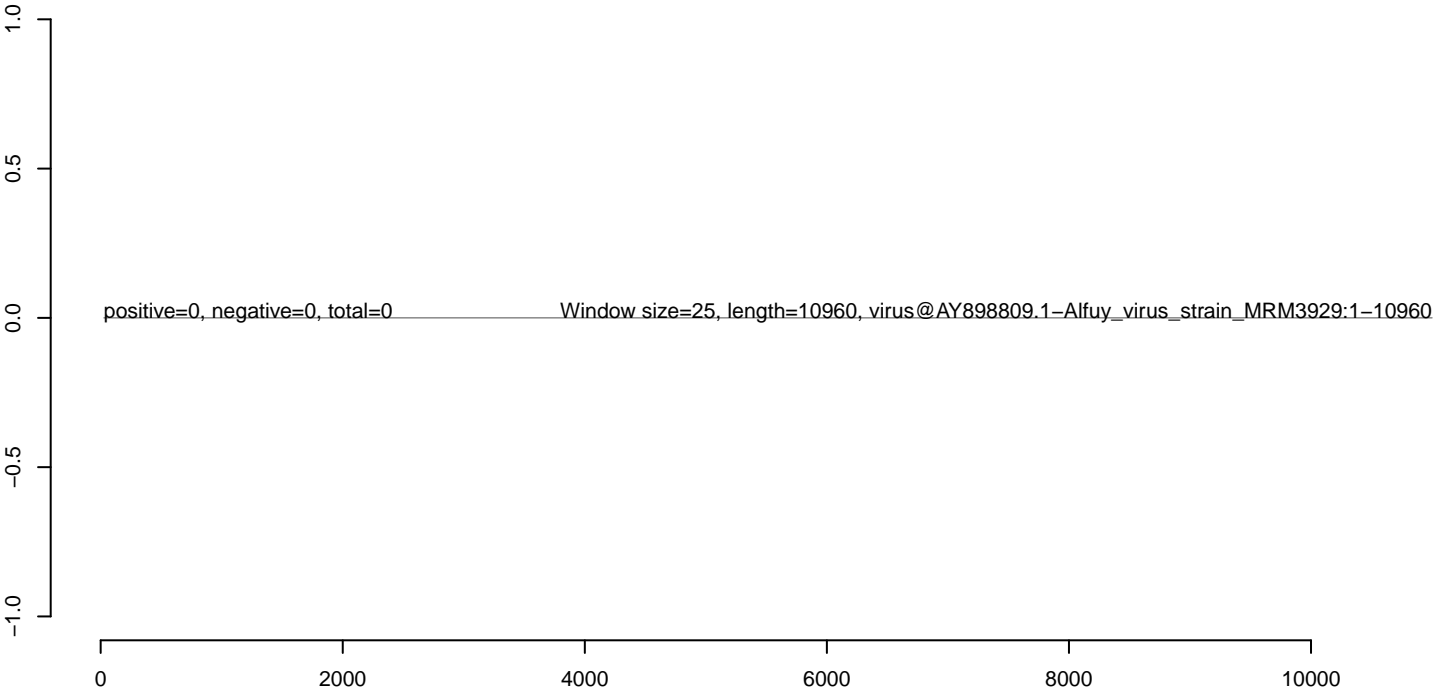
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



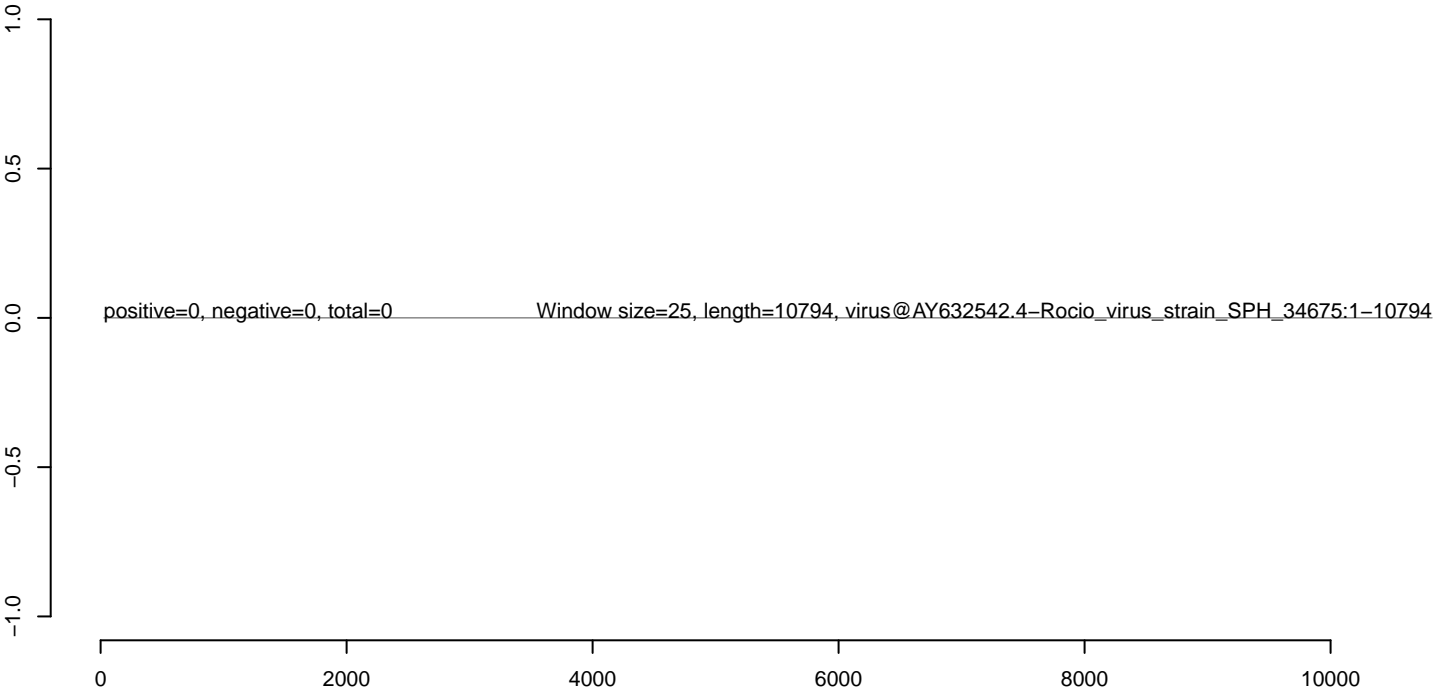
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



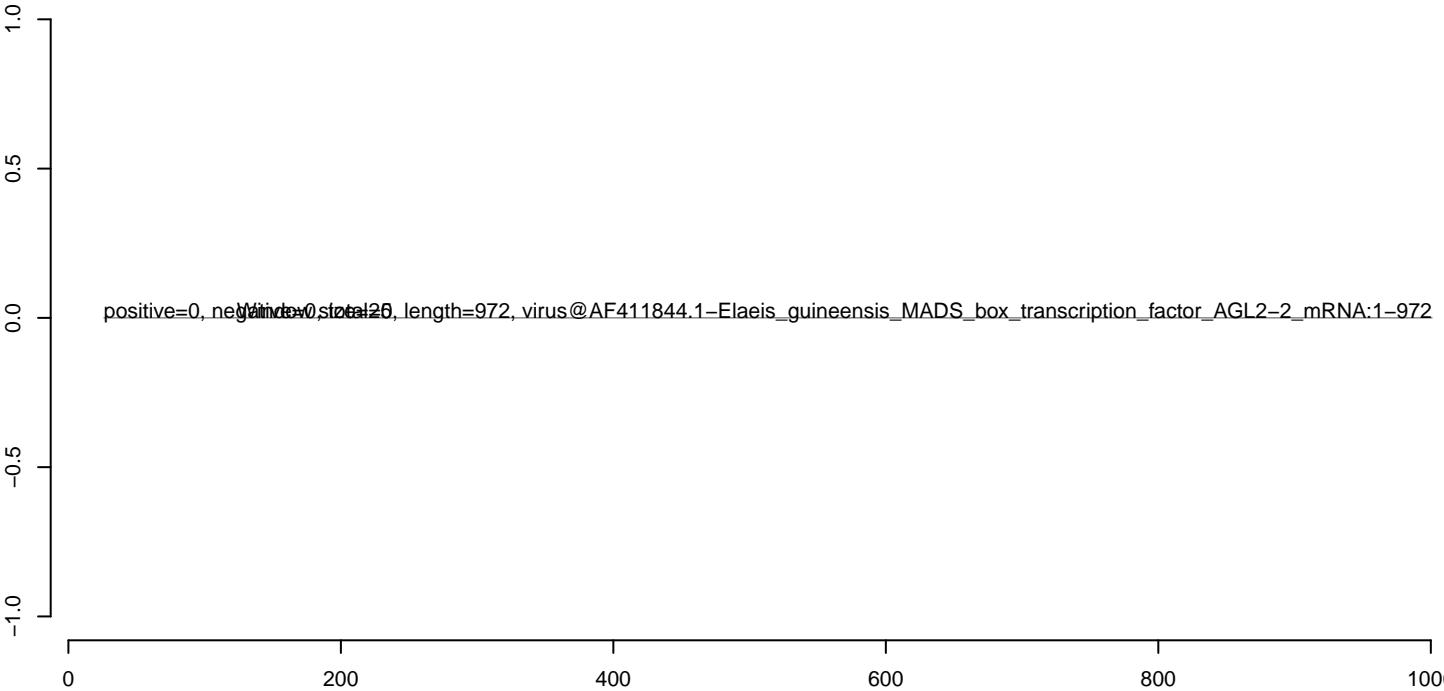
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



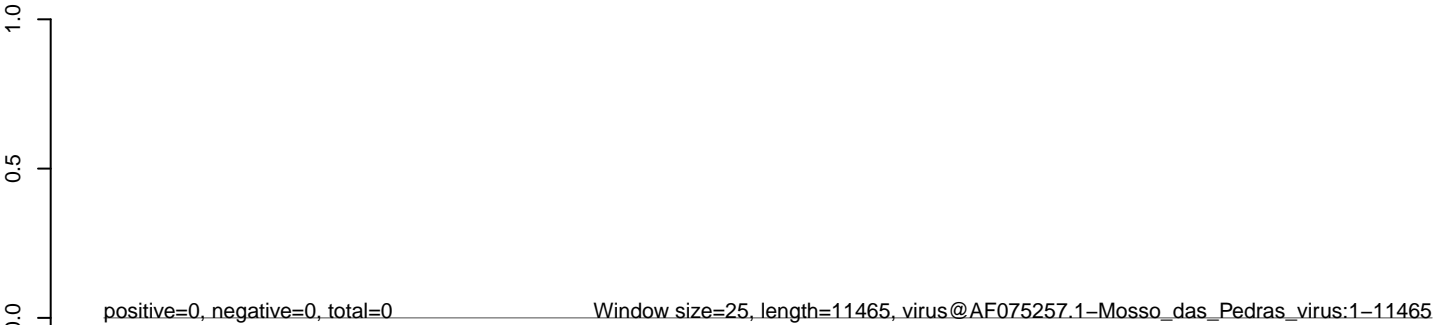
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



0 2000 4000 6000 8000 10000

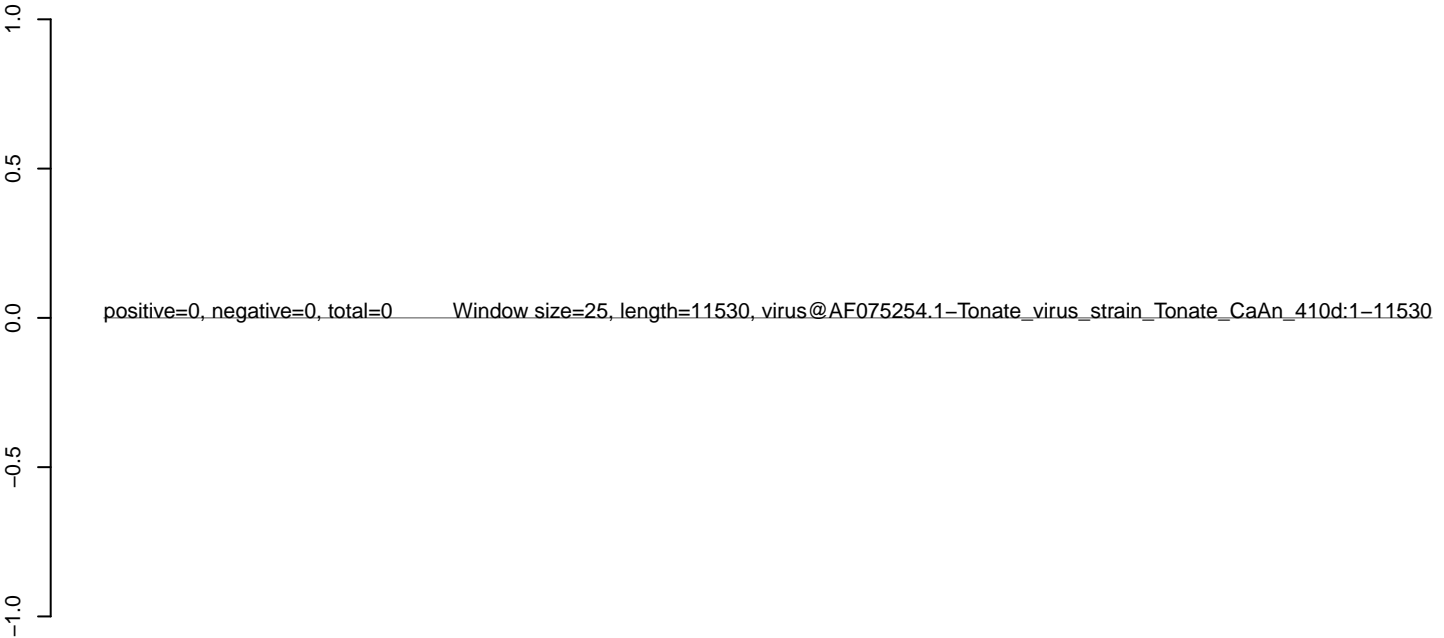
AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep



AeAlbo_C636_cells.18_23.rep



AeAlbo_C636_cells.24_35.rep



AeAlbo_C636_cells.rep

